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Microbial deglycosylation and ketonization of ginsenoside by *Cladosporium cladosporioide* and their anticancer activity

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Abstract Ginseng has been used for thousands of years in Asian countries as a traditional medicinal herb and has gained great popularity in the past decade. Ginsenosides are the major pharmacological components in ginseng. We here show that *Cladosporium cladosporioide* is able to convert the major ginsenoside Rb1 into four known metabolites (ginsenosides Rd, F2, CK and PPD) and two new metabolites [12 β -hydroxydammar-3-one-20(*S*)-*O*- β -D-glucopyranoside (3-oxo-CK) and dammar-24-en-12 β ,20(*S*)-diol-3-one (3-oxo-PPD)]. CK, PPD and 3-oxo-PPD were shown to have a potent antiproliferative activity against A549 lung cancer cells. We found that Rb1 \rightarrow Rd \rightarrow F2 \rightarrow CK \rightarrow PPD or 3-oxo-CK \rightarrow 3-oxo-PPD represents the ginsenoside metabolic pathway.

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Introduction

As a traditional medicinal herb, ginseng has been used for thousands of years in the Far East. It has gained great popularity in the West during the past decade (Attele et al. 1999; Ang-Lee et al. 2001). Ginsenosides are the major pharmacological components in ginseng with Rb1, Rb2, Rc, Rd, Rg1 and Re as the major compounds (Yang et al. 2014; Cheng et al. 2008). For instance, Rb1 is the major component of the protopanaxadiol group saponins in roots, making up 23.8 % of the total ginsenosides (Son et al. 2008).

Orally ingested major ginsenosides are activated by intestinal bacterial deglycosylation (Hasegawa 2004). In recent decades, many studies have reported the successful transformation of major ginsenosides into more active ginsenosides such as Rg3, CK and PPD. For instance, CK has anti-cancer (Kim et al. 2009; Ming et al. 2011), anti-angiogenic (Jeong et al. 2010), anti-inflammation (Joh et al. 2011) and hepatoprotective effects (Lee et al. 2005), while PPD provides antilung cancer (Zhang et al. 2013) and anti-prostate cancer activity (Cao et al. 2014). The natural availability of these ginsenosides in ginseng is low. The conversion of major ginsenosides to the more active minor ginsenosides can be accomplished through a number of methods such as acid treatment (Bae et al. 2004), alkali treatment (Yang et al. 2003), heating (Sun et al. 2009) and microbial transformation (Wu et al. 2012). Chemical transformation produces non-specific racemic mixtures of ginsenosides, which is why Microbial transformation is preferred (Quan et al. 2011). Microbial modification includes side-chain oxidation-reduction, hydroxylation and ketonization (Liu et al. 2011; Chen et al. 2013; Jin et al. 2014).

Lung cancer is the leading cause of cancer death in the world (Ferlay et al. 2010). Non-small-cell lung carcinoma (NSCLC) is the most common type of lung cancer accounting for 85–90 % of the cases (Gong et al. 2011). A large number of bioactive compounds are used to treat cancers, including vinblastine, paclitaxel and camptothecin. Still it is important to search for alternative therapeutic agents. Here we show that *Cladosporium cladosporioide* KACC 43926 converts Rb1 into several ginsenosides with potent antiproliferative activity against A549 lung cancer cells.

Materials and methods

Materials and organisms

Yeast mold (YM) broth was purchased from Difco (USA). The Silica gel 60 F_{254} plates and silica gel 60 (Merck, Germany) was used for TLC and column chromatography. All chemicals and solvents were analytical or HPLC grade. The strain *C. cladosporioide* KACC 43926 was purchased from Korean Agricultural Culture Collection (Suwon, Republic of Korea).

Preparation of ginsenoside Rb1

American ginseng extract with a ginsenoside content $\geq 80 \%$ (w/w) was purchased from Jiuhui Co. Ltd. (Changsha, Hunan, China). The crude extract was subjected to a silica gel column and eluted with CHCl₃-MeOH-H₂O (5:1:3–65:35:5, v/v/v). The eluate was then purified by semi-preparative HPLC to yield pure Rb1.

Biotransformation of ginsenoside Rb1

C. cladosporioide KACC 43926 was incubated in 150 mL YM broth containing 0.4 mg/mL Rb1 as the

carbon source in flasks at 28 °C and 150 rpm. After 10 days of incubation, the reaction mixture was extracted three times with water-saturated butanol. TLC analysis was performed using Silica Gel 60 plates and CHCl3-CH3OH-H2O (65:35:10 v/v/v). Compounds were detected by spraying 10 % (v/v) H_2SO_4 followed by heating at 110 °C for 10 min (Shibata et al. 1965). HPLC was performed using an Agilent 1260 system (Agilent). The separation was performed on a C18 column (50 \times 4.6 mm, ID 2.6 μ m) with H₂O (solvent A) and acetonitrile (solvent B) at A/B ratios of 81:19, 81:19, 71:29, 71:29, 60:40, 44:56, 30:70, 10:90, 10:90, 81:19, and 81:19; with run times of 0-7, 7-11, 11-14, 14-25, 25-28, 28-30, 30-31.5, 31.5-34, 34-34.5, and 34.5-40 min, respectively. The flow rate was 0.6 mL min⁻¹ and detection wavelength was 203 nm.

Isolation of metabolites

Fractions A–C were obtained by separating the reaction extract on silica gel column chromatography using CHCl₃–CH₃OH (13:1). Fraction A was separated with CH₂Cl₂–EtOH (50:1–1:1) yielding fractions D-E. Fraction were further purified by semipreparative HPLC. Metabolites were dissolved in pyridine- d_5 and identified by ¹H, ¹³C, and 2D NMR using an FT-NMR spectrometer (400 MHz; Varian Inova AS 400, Varian, Palo Alto, CA, USA). Chemical shifts are given in δ (ppm) based on tetramethyl-silane (TMS) as internal standard.

Proton NMR data of metabolite **5** were: ¹H-NMR (400 MHz, pyridine- d_5 , δ_H) 5.25 (1H, dd, J = 6.8, 6.4 Hz, H-24), 5.16 (1H, d, J = 7.6 Hz, H-1'), 4.46–3.89 (sugar moieties), 1.60 (9H, s, H-28, 21, 29), 1.11 (3H, H-26), 1.03 (3H, s, H-27), 0.96 (3H, s, H-18), 0.91 (3H, s, H-19), 0.87 (3H, s, H-30). Proton NMR data of metabolite **6** were: ¹H-NMR (400 MHz, pyridine- d_5 , δ_H) 5.47 (1H, dd, J = 6.4, 5.6 Hz, H-24), 3.89 (1H, m, H-12), 1.69 (3H, s, H-28), 1.66 (3H, s, H-21), 1.40 (3H, s, H-29), 1.13 (3H, H-26), 1.04 (3H, s, H-27), 0.99 (3H, s, H-18), 0.92 (3H, s, H-19), 0.87 (3H, s, H-30).

Cytotoxicity assay

Cell viability was measured by using the 3-(4,5dimethylthiazol-2-yl)-2,5-diphenyltetrazolium (MTT) assay (Mosmann 1983). A549 Lung Cancer cells were



Fig. 1 TLC analysis of metabolites of ginsenoside Rb1 converted by *C. cladosporioide* KACC 43926. C, ginsenoside Rb1 control, without inoculation of the strain; S, ginsenosides standard. The samples were withdrawn at different time. 1, 2 h; 2, 1 day; 3, 4 days; 4, 7 days; 5, 10 days

cultured for 24 h with different 0-300 μ M ginsenoside in 96-well plates at a density of 1 × 10⁴ cells/well. After incubation at 37 °C in a humidified incubator containing 5 % CO₂, 10 μ L of MTT (5 mg/mL) was added to each well and incubated for an additional 4 h. Precipitated formazan was dissolved in 100 μ L of DMSO for 30 min. The absorbance was recorded with a plate reader (Bio-Tek Instrument, USA) at a test wavelength of 570 nm and a reference wavelength of 630 nm.

Results

Biotranformation of ginsenoside Rb1

C. cladosporioide KACC 43926 converted ginsenoside Rb1 into several products as shown by TLC and HPLC analysis. Reaction products were separated on silica gel column chromatography. As a result, 4 known metabolites (ginsenoside Rd (metabolite 1), F2 (metabolite 2), CK (metabolite 3) and PPD (metabolite 4)) and 2 new metabolites (metabolites 5, 6) were obtained (Figs. 1, 2). Metabolite 1, 2 are intermediate metabolites, while metabolites 3, 4, 5, 6 represent final products.

Structure of metabolites

Metabolite **5** was obtained as a white powder. The ¹H and ¹³C NMR and DEPT (distortionless enhancement by polarization transfer) spectra of metabolite **5** were very similar to CK except for the appearance of a ketone (δ_C 216.2, C-3) and the disappearance of an oxygenated methine signal. The molecular weight of



Fig. 2 HPLC profiles of metabolites of ginsenoside Rb1 transformed by C. cladosporioide KACC 43926

 Table 1
 ¹³C-NMR spectrum of metabololites 5 and 6

Metabolite 5				Metabolite 6	
Aglycone moiety		Sugar moiety		Aglycone moiety	
No.	ppm	No.	ppm	No.	ppm
C-1	39.7	C-1′	98.2	C-1	39.6
C-2	34.2	C-2′	75.1	C-2	34.1
C-3	216.3	C-3′	79.2	C-3	215.9
C-4	47.3	C-4′	71.7	C-4	47.0
C-5	55.3	C-5′	78.2	C-5	55.0
C-6	19.9	C-6′	62.9	C-6	20.8
C-7	36.1			C-7	35.6
C-8	39.9			C-8	39.4
C-9	49.9			C-9	49.4
C-10	36.9			C-10	36.6
C-11	30.7			C-11	31.0
C-12	70.0			C-12	70.5
C-13	49.6			C-13	48.5
C-14	51.4			C-14	51.4
C-15	30.7			C-15	31.0
C-16	26.6			C-16	26.8
C-17	51.6			C-17	51.3
C-18	17.2			C-18	16.6
C-19	17.7			C-19	17.4
C-20	83.3			C-20	72.7
C-21	22.3			C-21	20.8
C-22	36.1			C-22	35.6
C-23	23.15			C-23	22.7
C-24	126.0			C-24	126.0
C-25	130.9			C-25	130.5
C-26	25.7			C-26	25.5
C-27	17.7			C-27	17.4
C-28	26.7			C-28	26.5
C-29	15.6			C-29	15.2
C-30	15.8			C-30	15.7

metabolite **5** was 2 daltons less than that of CK, indicating that the ketonization was introduced at C-3. The proton and carbon signals attributed to the sugar moiety suggested the presence of a β -glucopyranosyl group. The correlation between $\delta_H 5.16$ (H-1') and $\delta_C 83.3$ (C-20) in the HMBC (heteronuclear multiple bond correlation) spectrum and the chemical shift of the anomeric carbon signal ($\delta_C 98.2$) supported the presence of a glucopyranosyl group at C-20. Therefore, the structure of metabolite **5** was determined as

 12β -hydroxydammar-3-1-20(*S*)-*O*- β -D-glucopyranoside (3-oxo-CK).

Metabolite 6 was obtained as a white powder. The ¹³C-NMR spectrum of metabolite **6** displayed 30 carbon signals. All signals could be assigned based on a DEPT experiment, the HSQC (heteronuclear single quantum coherence) spectrum, and a comparison with the ¹³C-NMR data of PPD (Asakawa et al. 1977) (Table 1). The data were similar to PPD with the exception of the proton and carbon resonances indicating the presence of a carbonyl functional group instead of an oxygenated methane at the C-3 position. The ¹H and ¹³C NMR spectra (Table 1) showed signals of an olefine methine group [$\delta_{\rm H}$ 5.47 (J = 6.4, 5.6 Hz, H-24), δ_c 126.0 (C-24)], eight tertiary methyl groups (Table 1) and an oxygenated methine group $[\delta_H 3.89 (H-12), \delta_c 70.5 (C-12)]$. A ketone signal at δ_c 215.9 (C-3) was observed in the low magnetic field. In comparison to a previously isolated compound, metabolite 6 lacked a sugar moiety at the C-20 position. Consequently, the structure of metabolite 6 was determined to be dammar-24-en- 12β ,20(S)-diol-3-one (3-oxo-PPD) (Anufriev et al. 1997).

Biotransformation pathway

TLC analysis was performed to identify the metabolic pathway of the ginsenosides. To this end, samples of the reaction mixture were taken in time (Fig. 3). Ginsenoside Rb1 was converted into ginsenoside Rd by hydrolysis of a glucose unit at C-20 position. Then, ginsenoside F2 was produced from ginsenoside Rd by additional hydrolysis of a single glucose moiety at C-3 position. Ginsenoside F2 was converted into CK by hydrolysis of a glucose unit at C-3 position. Ginsenoside CK was transformed into PPD by hydrolysis of a glucose unit at C-20 position or 3-oxo-CK by ketonization at C-3 position. Finally, ginsenoside PPD and 3-oxo-CK were transformed into 3-oxo-PPD by ketonization at C-3 position and hydrolysis of a glucose unit at C-20 position, respectively. These results suggest that C. cladosporioide KACC 43926 has potent β -glucosidase and ginsenoside dehydrogenase activity.

In vitro cytotoxicity assay in A549 lung cancer cells

Cell viability effects of four derivatives (CK, 3-oxo-CK, PPD, 3-oxo-PPD) on A549 cells was evaluated by Fig. 3 Microbial transformation pathway of ginsenoside Rb1 by *C. cladosporioide* KACC 43926



Fig. 4 Effects of CK, PPD and 3-oxo-PPD on A549 cells viability. A549 cells were treated with different doses of CK, PPD, 3-oxo-PPD, and then measured by MTT assay. *p < 0.05, ****p < 0.001, ****p < 0.0001 for compounds-treated versus DMSO-treated cells. Mean values \pm SD (n = 3)



MTT assay. The 3-oxo-CK did not affect A549 cell growth in 24 h (data not shown). However, proliferation of A549 cells was suppressed in a dose-dependent way by CK, PPD and 3-oxo-PPD (Fig. 4). CK, PPD and 3-oxo-PPD significantly inhibited the growth of A549 cells at 50, 100 and 300 μ M, while 3-oxo-PPD only showed activity at 10 μ M. Thus, CK and PPD are more cytotoxic than 3-oxo-PPD.

Discussion

The natural major ginsenosides are excreted after uptake by the human body but when modified by the intestinal microbiota these bioactive compounds are retained. For instance, the ginsenosides Rb1, Rb2, Rc and Rd are metabolized into CK by intestinal microbiota (Hasegawa et al. 1996; Akao et al. 1998). Oral administration of ginsenoside Rb1 resulted in eight metabolites in rats, including gypenoside XVII, ginsenosides Rd, F2, CK and 3-oxo-CK (Chen et al. 2008). In addition, Jin et al. (2013) reported that PPD was metabolized into 3-oxo-PPD in rats after oral administration. It should be noted that individuals have their own characteristic indigenous gut microbiota (Simon and Gorbach 1986; Rumney and Rowland 1992; Wakabayashi et al. 1997) and their ginsenoside modifying activity can thus be different. Moreover, microbial intestinal composition can be affected disease, unbalanced diet, stress and lifestyles. Hence, the pharmacological effects of ginseng will depend on how effectively a person can metabolize and absorb ginsenosides. To circumvent this, ginsenosides can be modified in vitro.

This study shows for the first time that *C.* cladosporioide KACC 43926 can convert the major ginsenoside Rb1 into bioactive compounds that can also be found after oral administration. Evidence was presented that the metabolic pathway of Rb1 follows Rb1 \rightarrow Rd \rightarrow F2 \rightarrow CK \rightarrow PPD or 3-oxo-CK \rightarrow 3oxo-PPD. *C. cladosporioide* KACC 43926 has potent β -glucosidase for hydrolysis of a glucose unit at C-3 or C-20 position and ginsenoside dehydrogenase activity for ketonization at C-3 position.

The non-polar ginsenosides, such as Rg3, Rh2, Rk1, Rg5, CK and PPD are taken up in human breast cancer cells. The most non-polar ginsenoside PPD has the highest uptake rate, followed by CK (Ha et al. 2010). The Rb1 metabolites of *C. cladosporioide* should thus be easily absorbed in the body even in the absence of an intestinal microbiota that can metabolize ginsenosides. Based on the cytotoxicity results of four compounds, the ketonized compounds may reduce the cytotoxicity in cell lines. CK, PPD and 3-oxo-PPD significantly inhibited growth of A549 lung cancer cells at 50–300 μ M. Therefore, these compounds have both high uptake and potent antiproliferative activity against these cells.

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