Stereo and region-selective biosynthesis of two new dihydroartemisinic acid glycosides by suspension-cultured cells of *Artemisia annua*

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ABSTRACT

**Background:** The system of plant-cultured cells is one of the optimal systems to investigate biosynthesis pathway and their bioactive intermediates. **Objective:** To study the biosynthesis of dihydroartemisinic acid (1) by suspension-cultured cells of *Artemisia annua*. **Materials and Methods:** Substrate (compound 1) was administered into the suspension-cultured cells of *A. annua* and co-cultured for 2 days. The methanol extract was separated on various column chromatography methods and the structures of two biosynthesis products were elucidated based on the analysis of 1H NMR, 13C NMR, 2D NMR, and ESI-MS. Time-course curve was also established. Furthermore, in vitro antitumor activities of compounds 1-3 against HepG2, K562, and A549 cell lines were evaluated by MTT assay. **Results:** Two new compounds were obtained, namely 3α-hydroxy-dihydroartemisinic acid-α-D-glucopyranosyl ester (2) and 15-hydroxy-cadin-4-en-12-oic acid-β-D-glucopyranosyl ester (3). The results demonstrated that the cultured cells of *A. annua* possessed the abilities to stereo-selective hydroxylate and region-selective glycosylate sesquiterpene compounds in a highly efficient manner. Inhibitory effects of compounds 1-3 on proliferation of HepG2, K562, and A549 cell lines in vitro were also investigated. **Conclusion:** Two new dihydroartemisinic acid glycosides were obtained by stereo- and region-selective biosynthesis with cultured cells of *A. annua*. **Key words:** Antitumor, *Artemisia annua*, biosynthesis, dihydroartemisinic acid, glycosylation

INTRODUCTION

Artemisinin, a sesquiterpene lactone with a peroxide bridge extracted from Chinese medicinal herb (*Artemisia annua*, Qinghao), is famous for its bioactivity against both chloroquine-resistant and sensitive strains of *Plasmodium falciparum* as well as cerebral malaria with high safety profile.[1,2] In addition to their antimalarial activity, artemisinins were reported in recent decades as potential reagents against cancer cells.[3] Since artemisinin was found, its biosynthesis pathway attracted more and more attention. To use biotechnology method to produce or increase the yield of artemisin, many scientists have devoted themselves to elucidate the biosynthesis pathway.[4-8]

Dihydroartemisinic acid (1), one of the precursors of artemisinin,[7] attracted increasing attention after it was confirmed to transform to artemisinin *in vitro* by a nonenzymatic process.[7,8] Many researchers have been carried out to investigate the biosynthesis pathway from dihydroartemisinic acid to artemisinin.[5,6] However, the process is still incompletely understood, particularly within the plant cells.

Over the past few decades, biotransformation has been extensively studied because it is considered to be an important method for converting inexpensive and plentiful substances into expensive and scarce ones. Recently, plant cell cultures have been studied as potential agents during biotransformation reactions, especially for obtaining chiral alcohols, which are intermediates of pharmaceutical, and other potential compounds in industrial scale. Plant cell cultures as important biotransformation systems have been used widely.[9,12] To elucidate and evaluate the biosynthesis pathways of artemisinins, our research group has been screening tens of plant culture cell systems and has got some encouraging scientific information.[13-18]
To date, there is no report on the biotransformation of dihydroartemisinic acid by plant-cultured cells except the report from our research group. As a continuation work to explore the plant cell biotransformation of the precursor of artemisinin, and with the aim to enrich the metabolites of dihydroartemisinic acid, and to find out novel artemisinin derivatives, which might possess good antimalarial and/or antitumor activities, the biotransformation of dihydroartemisinic acid by suspension-cultured cells of *Artemisia annua* was investigated in the present paper.

**MATERIALS AND METHODS**

**General**

$^1$H and $^{13}$C nuclear magnetic resonance (NMR) and 2D NMR spectra were recorded on a Bruker DRX-400 spectrometer, the chemical shifts ($\delta$) were given in ppm relative to TMS as an internal standard, and coupling constants were given in Hz. ESI-MS data were obtained with a 4000 Q TRAP LC/MS/MS system by direct inlet using MeOH as solvent. HR-TOF-MS were recorded on SYNAPTEM G2 HDMS, Waters, Manchester, U.K. Silica gel (100-200 mesh and 200-300 mesh) used for column chromatography (CC), and silica GF$_2$54 (10-40 $\mu$m) for TLC were supplied by the Qingdao Marine Chemical Factory, China. ODS (YMC Co., Ltd., Japan) and Sephadex LH-20 (Pharmacia Co.) were also used for separation. HPLC analysis was performed on an Agilent 1200 liquid chromatography system (Palo Alto, CA, USA), equipped with vacuum degasser, quaternary gradient pump, auto-sampler, and DAD, connected to an Agilent ChemStation software. An Agilent Hypersil ODS column ($\phi$4.6 mm × 250 mm, 5 $\mu$m) was used. A binary gradient elution system consisted of water (A) and methanol (B) and separation was achieved using the following gradient program: 0–5 min 40–50% B; 5–10 min 50–60% B; 10–15 min 60–70% B; 15–20 min 70–85% B; 20–25 min 100% B, and finally, reconditioning the column with 40% B isocratic for 2 min. The flow rate was 0.8 ml/min, and the system operated at 30°C. The detection wavelength was set at 210 nm.

**Substrate**

Dihydroartemisinic acid (1) was extracted and isolated from *A. annua* by our research group according to the referenced protocol. The structure was determined by MS and NMR. Its purity was >98% by HPLC analysis.

**Plant cell cultures**

The cells of *A. annua* have been subcultured routinely every 3 weeks using MS medium containing 2,4-dichlorophenoxyacetic acid (2,4-D 0.5 mg/L) and 6-benzylaminopurine (6-BA 1 mg/L). Prior to being used for biotransformation experiments, the cultured cells were transferred to a 500-ml conical flask containing 200 ml medium and cultured on a rotary shaker (110 rpm) for 13 days at 25°C in the dark.

**Biotransformation of dihydroartemisinic acid (1)**

Dihydroartemisinic acid (1, 110 mg) was dissolved in 1.1 ml of ethanol and distributed among 22 Erlemeyer flasks of 13-day-old cultures and incubated for an additional 2 days. After incubation, the cultures and media were separated by filtration with suction. The dried cultures were extracted with methanol for four times by ultrasound-assisted extraction. Each of the MeOH fractions was concentrated and partitioned between H$_2$O and EtOAc. The EtOAc fractions were combined and further purified on column chromatography by silica gel, sephadex LH-20, and ODS to afford products 2 and 3. The same culture was repeated 5 times.

**3 α-Hydroxydihydroartemisinic acid-α-D-glucopyranosyl ester (2)**

1H NMR (pyridine-d$_6$, 400 MHz) $\delta$: 0.8 (3H, d, $J = 4.4$ Hz), 1.2 (3H, d, $J = 6.8$ Hz), 2.0 (3H, s), 2.61 (1H, br, s), 5.31 (1H, s), 6.32 (1H, d, $J = 3.6$ Hz); $^{13}$C-NMR (pyridine-d$_6$, 100 MHz) $\delta$: 42.4 (C-1), 35.2 (C-2), 67.0 (C-3), 139.6 (C-4), 121.7 (C-5), 37.1 (C-6), 44.6 (C-7), 27.4 (C-8), 37.0 (C-9), 28.8 (C-10), 43.7 (C-11), 175.9 (C-12), 14.9 (C-13), 19.6 (C-14), 20.2 (C-15), 95.6 (C-1′), 73.9 (C-2′), 78.3 (C-3′), 70.8 (C-4′), 79.1 (C-5′), 61.9 (C-6′); ESI-MS: $m/z$ 413 [M-H]$^-$, 437 [M+Na]$^+$, calcd. for C$_{21}$H$_{34}$O$_4$Na$,^+$, 437.2151.

**15-Hydroxy-cadin-4-en-12-oic acid-β-D-glucopyranosyl ester (3)**

1H NMR (pyridine-d$_6$, 400 MHz) $\delta$: 0.744 (3H, d, $J = 6.4$ Hz), 1.21 (3H, d, $J = 6.8$ Hz), 1.79 (3H, s), 2.57 (1H, br, s), 4.30 (2H, s), 5.89 (1H, s), 6.32 (1H, d, $J = 8.0$ Hz); $^{13}$C-NMR (pyridine-d$_6$, 100 MHz) $\delta$: 43.5 (C-1), 27.0 (C-2), 23.8 (C-3), 142.1 (C-4), 120.7 (C-5), 37.9 (C-6), 45.2 (C-7), 28.9 (C-8), 36.7 (C-9), 28.9 (C-10), 43.9 (C-11), 177.4 (C-12), 16.5 (C-13), 21.0 (C-14), 67.9 (C-15), 97.1 (C-1′), 75.4 (C-2′), 79.9 (C-3′), 72.3 (C-4′), 80.6 (C-5′), 63.5 (C-6′); ESI-MS: $m/z$ 413 [M-H], 437 [M+Na]$^+$, HR-TOF-MS ($m/z = 437.2144$ [M + Na]$^+$), calcd. for C$_{21}$H$_{34}$O$_4$Na$,^+$, 437.2151.

**Time-course of biotransformation**

Cultured cells of *A. annua* (10 g) were transferred to a 500-ml Erlemeyer flask containing 200 ml medium, and cultured by continuous shaking for 13 days at 25°C. Compound 1 (5 mg/flask) was added to the suspension cultures and incubated at 25°C in a rotary shaker (110 rpm). At 1-day intervals, three of the flasks were taken out from the culture to extract the samples for HPLC analysis. The samples were then separated and the HPLC chromatograms were compared with those of the standard samples to determine the concentration of each compound.

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filtration. The extraction and analysis procedures were the same as those described earlier. The yields of the products were calculated on the basis of the peak area from HPLC using calibration curves prepared by HPLC-DAD and were expressed as relative percentages to the total amount of whole biosynthesis products.

**MTT-cell proliferation assay**

The inhibitory effects of 1-3 on the proliferation of HepG2, K562, and A549 cells were evaluated in vitro by MTT [3-(4,5-dimethylthiazole-2-yl)-2,5-diphenyltetrazoliumbromide] assay, which was performed as described in the literature. The concentrations of biosynthesis products on the selected cell lines were in the range of 0.0185-0.5000 μmol/ml.

**RESULTS**

**Structural elucidation of biotransformation products**

Two new compounds were obtained after dihydroartemisinic acid (1) was incubated with the plant-cultured cells of A. annua for 2 days. Their structures were elucidated to be 3α-hydroxy-dihydroartemisinic acid-α-d-glucopyranosyl ester (2) and 15-hydroxy-cadin-4-en-12-oic acid-β-d-glucopyranosyl ester (3) [Figure 1].

Compound 2 was obtained as an amorphous powder. It displayed a quasi-molecular ion peak at m/z 413 [M-H]- and 437 [M+Na]+ in ESI-MS, indicating that the molecular weight of product 2 was 414, 178 more than that of dihydroartemisinic acid. The molecular formula of 2 was determined as C21H34O8 on the basis of HR-TOF-MS (m/z = 437.2154 [M+Na]+, calcd. for C21H34O8Na+, 437.2151). This information, together with the 13C NMR spectrum, suggested that compound 2 might be a molecule that has a glucose attached to hydroxyl-dihydroartemisinic acid. In the 'H NMR spectrum of 2, in addition to the signals for dihydroartemisinic acid moiety, it showed one set of α-glucopyranosyl signals, with the resonance for the anomeric proton at δ 6.32 (1H, d, J = 3.6 Hz). Its 13C NMR spectrum exhibited 21 carbon signals, including one anomeric carbon signal at δ 95.6, suggesting that 2 was larger than 1 by one hexose moiety and a hydroxyl group. A comparison of 13C NMR spectra of 2 with that of 1 showed that carboxyl carbon signal was shifted upfield by 7.9 ppm (183.8 → 175.9). This suggested that 2 was an ester. The carbon signal at C-15 in 2 was shifted upfield (δ 23.7 → 20.2), and one carbon signal was shifted markedly downfield (δ 26.6 → 67.0). These data suggested that the upfield shift of the C-15 signal was caused by a γ-effect due to hydroxylation of 1 at the position of C-3. The 13C NMR signal of the aglycone in 3 was coincident with that of 3α-hydroxy-dihydroartemisinic acid.[9] HMBC spectrum showed that δ 175.9 (C12) was correlated with δH 6.32 (1H, d, J = 3.6 Hz), suggesting that sugar moiety was linked to carbonyl group of 1. Therefore, the structure of 2 was proposed to be 3α-hydroxy-dihydroartemisinic acid-α-d-glucopyranosyl ester. Biotransformation product 2 is a new compound.

Compound 3 possessed the same molecular weight as that of 2. The molecular formula of 3 was determined as C21H34O8 on the basis of HR-TOF-MS (m/z = 437.2144 [M + Na]+, calcd. for C21H34O8Na+, 437.2151). The main difference...
establishment of time-course curve of substrate 1

Figure 2 showed the results of the biotransformation products of compound 1 by cultured cells of *A. annua*. As indicated in [Figure 2], compound 1 disappeared after 3 days’ culture, meaning that 1 was completely transformed. A possible biosynthesis pathway in the cultures was proposed in [Figure 1]. As shown in [Figure 1], hydroxylation firstly happened at C-15 and C-4. Therefore, 3 was determined to be 15-hydroxy-cadin-4-en-12-oic-β-D-glucopyranosyl ester. Biotransformation product 3 is also a new compound.

** Establishment of time-course curve of substrate 1 **

**Table 1: Inhibitory effects of compounds 1-3 (0.5 µmol for each) on proliferation of HepG2, K562, and A549 cell lines**

| Cell lines | Inhibition rate (%) |
|------------|---------------------|
|            | 1                   | 2                   | 3                   |
| HepG2      | 37.4±3.7**          | 78.0±1.1**          | 70.7±1.5**          |
| K562       | 50.6±3.9**          | 25.4±1.6**          | 7.7±2.5             |
| A549       | 99.9±0.1**          | 16.9±2.9**          | 0±4.8               |

Values are given as means±standard deviation of three separate experiments. **Compared to control: *P<0.01***

**In vitro antitumor activities**

Antitumor activities of 1-3 against the HepG2, K562, and A549 cell lines are shown in [Table 1]. The sensitivity of those three cell lines to compounds 1-3 was quite different. For HepG2 cell line, the inhibitory rate of biotransformation products (2 and 3) was higher than that of compound 1. K562 cell line was not sensitive to both the substrate and the products. Compound 1 showed good inhibitory activity against the A549 cell line while no inhibitory activity appeared for the biotransformation products.

**DISCUSSION**

The experiment of time-course curve revealed that the yields of compounds 2 and 3 were not high enough while compound 1 was completely transformed. Those might due to the metabolites of dihydroartemisinic acid in cultured cells of *A. annua* were a wide variety. In our previous study, hydroxylation products and artemisinins were isolated as the metabolites of dihydroartemisinic acid by *A. annua*.

Therefore, the metabolism of dihydroartemisinic acid in cultured cells, such as *A. annua*, was very complicated and many metabolites might have existed.

There may be two metabolic pathways for 1 in cells of *A. annua*: To be biosynthesized to complex compounds, such as artemisinin, and to be decomposed to nontoxic constituents. Hydroxylation and glycosylation are supposed to be initial steps to biodegradation of toxicants in plants.

In our previous study, 15-hydroxy-cadin-4-en-12-oic acid was isolated as the metabolite of dihydroartemisinic acid when using crown galls of *Panax quinquefolium* as the biocatalyst system. At the present study, its ester was obtained. This demonstrated that both plant-cultured cells of *A. annua* and crown galls of *P. quinquefolium* could hydroxylate dihydroartemisinic acid at C-15 position.

More importantly, the glycosyltransferase of plant-cultured cells of *A. annua* selectively glycosylated dihydroartemisinic acid with carboxyl group at C-12 position to produce corresponding glycosides though hydroxyl group existed. This information indicated that the glycosyltransferase functioned as a high region-selective enzyme. In addition, α-glycoside was isolated in this experiment. Generally,
α-configuration compound is rarely found when compared with β-configuration products isolated from plant biotransformation system.[22] β-Glycosidase had been isolated from A. annua 24 years ago.[23] But the information about α-glycosidases of A. annua has not been reported up to now. This is the first report that the administrated substrate was converted into its d-glucopyranoside of α-configuration by cultured cells of A. annua. Characterization of enzyme that catalyzes the α-glycosylation is now in progress.

CONCLUSION

In summary, plant-cultured cells of A. annua have the ability to stereo-selective hydroxylate and region-selective glycosylate exogenous compounds like sesquiterpene in a highly efficient manner. The two new compounds (2 and 3) demonstrated excellent inhibitory effect on proliferation of HepG2 cell line in vitro.

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