

Biosynthesis of acteoside in cultured cells of *Olea europaea*

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Abstract Five phenylethanoid glycosides (acteoside, isoacteoside, β -oxoacteoside, β -hydroxyacteoside, and salidroside) were isolated from a cell suspension culture of *Olea europaea*. We examined the biosynthesis of acteoside in olive cell cultures by using feeding experiments with stable isotope labeled precursors. The hydroxytyrosol moiety of acteoside is biosynthesized from tyrosine through dopamine, whereas the caffeoyl moiety of acteoside is biosynthesized from phenylalanine via a cinnamate pathway. Dopamine is incorporated into acteoside through oxidation to the corresponding aldehyde, reduction to the alcohol, and then β -glycosylation.

Keywords Secondary metabolites · Olive · Verbascoside · 2-Phenethyl β -primeveroside · Hydroxytyrosol

Introduction

Olea europaea L. (olive) is an evergreen tree native to the Mediterranean coast whose fruit and oil have been used for food and cooking. There are reports about the olive's effectiveness in the prevention of heart ailments and hypertension [1]. The olive tree contains oleanane-type triterpenes, e.g., oleanolic acid, as a secondary metabolite;

the triterpenes are accumulated in the fruit [2]. We induced olive cell cultures and investigated their components, expecting to obtain a large amount of the oleanane-type triterpenes and triterpene glycosides; however, no triterpene glycosides were isolated, and more ursane-type triterpenes than oleanane-type triterpenes were isolated from the cell suspension cultures [3]. We also determined that the main secondary metabolites in olive cell cultures were phenylethanoid glycosides.

Acteoside (verbascoside) is a typical phenylethanoid glycoside, which has antioxidant [4], anti-inflammatory [5], antihypertensive [6], and various other biological and pharmaceutical uses. Acteoside has a 3,4-dihydroxyphenylethanol (hydroxytyrosol) moiety as a partial structure, which is comparatively rare in secondary metabolites [7–9]; however, the biosynthesis of acteoside and other phenylethanoid glycosides has not been thoroughly investigated [10, 11]. In this paper, we report biosynthetic studies of phenylethanoid glycosides by using experiments feeding stable isotope labeled compounds into olive cell cultures.

Materials and methods

General procedures

L-[1-¹³C]Phenylalanine (99% isotopic abundance), L-[3-¹³C]phenylalanine (99%), and L-[1-¹³C]tyrosine (99%) were purchased from Sigma-Aldrich Chemical Co. L-[3-¹³C]Tyrosine (99%) and [1-¹³C]phenethylamine (99%) were purchased from ISOTEC Inc. [2,2-D₂]Dopamine (97–98%), [1,1,2,2-D₄]dopamine (97–98%), L-[2,3-¹³C₂-4-hydroxy-¹⁸O]DOPA (97% [¹³C], 95% [¹⁸O]), and [1-¹³C]2-phenylethanol (99%) were purchased from

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Cambridge Isotope Laboratories Inc. Diaion HP20 was purchased from Mitsubishi Chemical Co. Millex-GP 0.22 μm was purchased from Millipore Co. Other general materials were purchased from Sigma-Aldrich Co., Merck Co., or Wako Pure Chemical Industries, Ltd. NMR spectra were recorded on a JEOL JNM-LA500 spectrometer in CD_3OD at 30°C . ^1H NMR spectra were recorded at 500 MHz and the chemical shifts were referenced to 3.30 ppm for the central peak of CHD_2OD in CD_3OD . ^{13}C NMR spectra were recorded at 125 MHz and the chemical shifts were referenced to 49.8 ppm for the central peak of CD_3OD . The relative ^{13}C abundance of individual carbon atoms (% ^{13}C) was calculated by comparing the ^{13}C signal integral between the ^{13}C -labeled product and non-labeled samples. The values were referenced to 1.1% for the carbon signal of glucose-1 observed at 104.2 ppm (see Table 1) for labeled and non-labeled samples. HPLC was performed on a Shimadzu model LC-10 system with a SPD-10Avp photodiode array detector and an RID-10A refractive index detector. LC–electron spray ionization (ESI)–MS was performed on a Thermoquest model LCQ system in negative mode, equipped with a Hewlett-Packard HP1100 system. LC was performed using a Shim-pack CLC-ODS(M) column (4.6×250 mm) with 33% MeOH (flow rate 1.0 ml/min) at 40°C . Relative MS peak intensity of each sample was referenced to 100 for the peak of m/z 623 [$\text{M} - \text{H}$] $^-$, the base peak observed with the natural acteoside.

Callus induction of *O. europaea* and subcultures

The callus induction of *O. europaea* into CM-NH_4 medium has already been reported [3]. The CM-NH_4 agar medium was a modified Murashige and Skoog's [12] agar medium with 30 g/l sucrose, 1 mg/l 2,4-dichlorophenoxyacetic acid, 0.1 mg/l kinetin, 7% coconut milk, and 9 g/l agar, supplemented with K^+ instead of NH_4^+ . The calli cultured on the CM-NH_4 agar medium were transferred to a DK-NH_4 agar medium, which was CM-NH_4 agar medium without coconut milk, and subcultured every 4 weeks at 25°C in the dark.

Extraction and isolation of phenylethanoid glycosides from the cultured cells of *O. europaea*

The olive cells (about 9 g, fresh weight (f.w.)) cultured for 4 weeks on a DK-NH_4 agar medium were inoculated into 200-ml conical flasks containing 80 ml of DK-NH_4 liquid medium and cultured on a rotary shaker (60 rpm) at 25°C in the dark. After four additional weeks of culturing, the cells were harvested and fresh cells from 6 flasks (190 g, f.w.) were homogenized and extracted twice with MeOH at room temperature. The MeOH extract solution was

Table 1 ^{13}C NMR (CD_3OD , 125 MHz) chemical shifts of compounds 1–5

Carbon position ^a	1	2	3	4	5
1	131.5	131.4	127.6	133.7	130.8
2	117.1	117.1	115.5	114.6	130.9
3	146.1	146.1	152.7	146.3	116.1
4	144.7	144.7	146.9	146.1	156.8
5	116.5	116.6	116.1	116.2	116.1
6	121.3	121.3	123.0	119.0	130.9
7	36.5	36.7	196.2	74.2	36.4
8	72.2	72.4	72.1	76.4	72.1
Glucose-1	104.2	104.4	104.1	104.6	104.4
Glucose-2	76.2	75.7	76.3	76.7	75.1
Glucose-3	81.7	84.0	81.2	81.3	78.1
Glucose-4	70.4	70.4	70.4	70.4	71.7
Glucose-5	76.0	75.4	76.4	76.1	77.9
Glucose-6	62.4	64.6	62.5	62.3	62.8
1'	127.7	127.6	127.5	127.6	
2'	115.3	115.1	115.2	115.2	
3'	148.0	146.8	146.9	146.9	
4'	149.8	149.7	149.9	149.9	
5'	116.3	116.4	116.5	116.5	
6'	123.2	123.1	123.2	123.2	
7'	146.8	147.3	148.1	148.0	
8'	116.5	114.8	114.6	114.6	
9'	168.3	169.1	168.2	168.3	
Rhamnose-1	103.0	102.7	103.0	102.9	
Rhamnose-2	72.2	72.3	72.1	72.1	
Rhamnose-3	72.1	72.4	72.1	72.4	
Rhamnose-4	73.8	74.0	73.8	73.8	
Rhamnose-5	70.6	70.1	70.4	70.4	
Rhamnose-6	18.4	17.9	18.4	18.4	

^a See Fig. 1 for numbering

concentrated. The extract was partitioned between hexane and H_2O . The H_2O layer was passed through a Diaion HP20 column. The column was washed with H_2O and then eluted with MeOH. The MeOH eluate was evaporated and then partitioned between *n*-BuOH and H_2O . The *n*-BuOH extract (151 mg) was subjected to HPLC repeatedly using Millipore Prep Nova-Pak HR columns (6 μm , C_{18} , 60 \AA , 7.8×300 mm), 32% MeOH, 2.0 ml/min to yield compounds **1** (104 mg), **2** (5 mg), **3** (3 mg), and **4** (12 mg). The eluate before compound **1** was concentrated and subjected to HPLC using the same column (20% MeOH, 2.0 ml/min) to yield compound **5** (2 mg).

Acteoside (**1**)

Amorphous; negative ESI MS m/z : 623 [$\text{M} - \text{H}$] $^-$; ^1H NMR δ : 1.05 (3H, d, $J = 6.5$ Hz, rhamnose-6), 2.78 (2H,

dd, $J = 8.0, 7.0$ Hz, H-7), 3.28 (1H, dd, $J = 9.5, 9.5$ Hz, rhamnose-4), 3.38 (1H, dd, $J = 9.5, 8.5$ Hz, glucose-2), 3.50–3.64 (5H, m, glucose-5, 6a, 6b, rhamnose-3, 5), 3.72 (1H, ddd, $J = 9.0, 8.0, 7.0$ Hz, H-8a), 3.81 (1H, dd, $J = 9.5, 9.5$ Hz, glucose-3), 3.91 (1H, dd, $J = 3.5, 1.5$ Hz, rhamnose-2), 4.04 (1H, ddd, $J = 9.0, 8.0, 7.0$ Hz, H-8b), 4.37 (1H, d, $J = 8.5$ Hz, glucose-1), 4.91 (1H, dd, $J = 9.5, 9.5$ Hz, glucose-4), 5.18 (1H, d, $J = 1.5$ Hz, rhamnose-1), 6.27 (1H, d, $J = 16.0$ Hz, H-8'), 6.56 (1H, dd, $J = 8.5, 2.0$ Hz, H-6), 6.67 (1H, d, $J = 8.5$ Hz, H-5), 6.69 (1H, d, $J = 2.0$ Hz, H-2), 6.77 (1H, d, $J = 8.5$ Hz, H-5'), 6.95 (1H, dd, $J = 8.5, 2.0$ Hz, H-6'), 7.05 (1H, d, $J = 2.0$ Hz, H-2'), 7.59 (1H, d, $J = 16.0$ Hz, H-7'); ^{13}C NMR (see Table 1).

Isoacteoside (2)

Amorphous; negative ESI MS m/z : 623 $[\text{M} - \text{H}]^-$; ^1H NMR δ : 1.24 (3H, d, $J = 6.0$ Hz, rhamnose-6), 2.78 (2H, ddd, $J = 7.0, 7.0, 2.0$ Hz, H-7), 3.28–3.34 (1H, m, glucose-2), 3.40 (1H, dd, $J = 9.0, 9.0$ Hz, glucose-4), 3.50–3.56 (2H, m, H-8a, glucose-5), 3.52 (1H, dd, $J = 9.0, 9.0$ Hz, glucose-3), 3.68–3.78 (1H, m, rhamnose-3), 3.82–4.00 (3H, m, H-8b, rhamnose-2, 4), 4.32 (1H, d, $J = 8.0$ Hz, glucose-1), 4.34 (1H, dd, $J = 12.0, 5.5$ Hz, glucose-6b), 4.49 (1H, dd, $J = 12.0, 2.0$ Hz, glucose-6a), 5.16 (1H, d, $J = 1.5$ Hz, rhamnose-1), 6.28 (1H, d, $J = 16.0$ Hz, H-8'), 6.52 (1H, dd, $J = 8.0, 2.0$ Hz, H-6), 6.63 (1H, d, $J = 8.0$ Hz, H-5), 6.66 (1H, d, $J = 2.0$ Hz, H-2), 6.76 (1H, d, $J = 7.5$ Hz, H-5'), 6.88 (1H, dd, $J = 7.5, 2.0$ Hz, H-6'), 7.03 (1H, d, $J = 2.0$ Hz, H-2'), 7.55 (1H, d, $J = 16.0$ Hz, H-7'); ^{13}C NMR (see Table 1).

β -Oxoacteoside (3)

Amorphous; negative ESI MS m/z : 637 $[\text{M} - \text{H}]^-$; ^1H NMR δ : 1.09 (3H, s, $J = 6.5$ Hz, rhamnose-6), 3.27–3.34 (1H, m, rhamnose-4), 3.50–3.65 (6H, m, glucose-2, 5, 6a, 6b, rhamnose-3, 5), 3.86 (1H, dd, $J = 9.5, 9.5$ Hz, glucose-3), 3.93 (1H, dd, $J = 3.0, 1.5$ Hz, rhamnose-2), 4.47 (1H, d, $J = 8.0$ Hz, glucose-1), 4.92 (1H, d, $J = 17.0$ Hz, H-8a), 4.93 (1H, dd, $J = 9.5, 9.5$ Hz, glucose-4), 5.22 (1H, d, $J = 1.5$ Hz, rhamnose-1), 5.26 (1H, d, $J = 17.0$ Hz, H-8b), 6.27 (1H, d, $J = 16.0$ Hz, H-8'), 6.77 (1H, d, $J = 8.5$ Hz, H-5), 6.82 (1H, d, $J = 8.0$ Hz, H-5'), 6.95 (1H, dd, $J = 8.0, 2.0$ Hz, H-6'), 7.05 (1H, d, $J = 2.0$ Hz, H-2'), 7.42 (1H, d, $J = 1.5$ Hz, H-2), 7.43 (1H, dd, $J = 8.5, 1.5$ Hz, H-6), 7.59 (1H, d, $J = 16.0$ Hz, H-7'); ^{13}C NMR (see Table 1).

β -Hydroxyacteoside (4)

Amorphous; negative ESI MS m/z : 639 $[\text{M} - \text{H}]^-$; ^1H NMR δ : 1.09 (3H, d, $J = 6.5$ Hz, rhamnose-6), 3.31 (1H, m, rhamnose-4), 3.45 (1H, dd, $J = 9.0, 8.5$ Hz, H-8a),

3.51–3.62 (6H, m, glucose-2, 5, 6a, 6b, rhamnose-2, 5), 3.84 (1H, dd, $J = 9.5, 9.5$ Hz, glucose-3), 3.92 (1H, dd, $J = 3.5, 2.5$ Hz, rhamnose-3), 3.97 (1H, dd, $J = 9.0, 3.0$ Hz, H-8b), 4.41 (1H, d, $J = 8.0$ Hz, glucose-1), 4.74 (1H, dd, $J = 8.5, 3.0$ Hz, H-7), 4.93 (1H, dd, $J = 9.5, 9.5$ Hz, glucose-4), 5.21 (1H, d, $J = 1.0$ Hz, rhamnose-1), 6.26 (1H, d, $J = 15.5$ Hz, H-8'), 6.70 (1H, d, $J = 8.5, 2.0$ Hz, H-6), 6.73 (1H, d, $J = 8.0$ Hz, H-5'), 6.77 (1H, dd, $J = 8.5$ Hz, H-5), 6.83 (1H, d, $J = 2.0$ Hz, H-2), 6.95 (1H, dd, $J = 8.0, 2.5$ Hz, H-6'), 7.04 (1H, d, $J = 2.5$ Hz, H-2'), 7.58 (1H, d, $J = 15.5$ Hz, H-7'); ^{13}C NMR (see Table 1).

Salidroside (5)

Amorphous; negative ESI MS m/z : 299 $[\text{M} - \text{H}]^-$; ^1H NMR δ : 2.82 (2H, dd, $J = 8.0, 6.5$ Hz, H-7), 3.17 (1H, dd, $J = 9.5, 7.5$ Hz, glucose-2), 3.25 (1H, ddd, $J = 9.5, 5.5, 2.5$ Hz, glucose-5), 3.26 (1H, dd, $J = 9.5, 9.0$ Hz, glucose-4), 3.34 (1H, dd, $J = 9.5, 9.0$ Hz, glucose-3), 3.65 (1H, dd, $J = 11.5, 5.5$ Hz, glucose-6b), 3.69 (1H, ddd, $J = 9.5, 8.0, 6.5$ Hz, H-8a), 3.85 (1H, dd, $J = 11.5, 2.5$ Hz, glucose-6a), 4.02 (1H, ddd, $J = 9.5, 8.0, 6.5$ Hz, H-8b), 4.28 (1H, d, $J = 7.5$ Hz, glucose-1), 6.68 (2H, d, $J = 9.0$ Hz, H-3,5), 7.05 (2H, d, $J = 9.0$ Hz, H-2,6); ^{13}C NMR (see Table 1).

Preparation of $[2-^{13}\text{C}]$ tyramine

$[2-^{13}\text{C}]$ Tyramine was prepared according to Sugimoto and Yamada [13]. A mixture of 100 mg L- $[3-^{13}\text{C}]$ tyrosine, 23 mg L-tyrosine decarboxylase (10 U), 25 ml 0.2 M acetate buffer (pH 5.5), and 50 ml H_2O was incubated for 8 h at 37°C. The incubation mixture was made alkaline (pH 10) with ammonia water and extracted with EtOAc. The EtOAc layer was evaporated and 42 mg $[2-^{13}\text{C}]$ tyramine was obtained.

Feeding experiments using stable isotope labeled compounds

The olive cells (about 9 g, f.w.) cultured for 4 weeks on a DK-NH₄ agar medium were inoculated into 200-ml conical flasks containing 80 ml of DK-NH₄ liquid medium and cultured on a rotary shaker (60 rpm) at 25°C in the dark. After an additional 3 weeks of culture, 10 mg L-phenylalanine ($[1-^{13}\text{C}]:[3-^{13}\text{C}] = 1:1$), 10 mg L-tyrosine ($[1-^{13}\text{C}]:[3-^{13}\text{C}] = 1:1$), 5 mg L- $[2,3-^{13}\text{C}_2-4\text{-hydroxy-}^{18}\text{O}]$ DOPA, 10 mg $[2,2\text{-D}_2]$ dopamine, 10 mg $[1,1,2,2\text{-D}_4]$ dopamine, 10 mg $[1-^{13}\text{C}]$ 2-phenylethanol, 10 mg $[1-^{13}\text{C}]$ phenethylamine, or 10 mg $[2-^{13}\text{C}]$ tyramine was administered to a cell suspension culture. The deuterium- or ^{13}C -labeled compounds were dissolved in 3–4 ml water and sterilized

by filtration (Millex-GP 0.22 μm). After administration, each culture was incubated for an additional week. Acteoside (**1**) was partially isolated (about 5 mg from each culture) as described above. In the $[2\text{-}^{13}\text{C}]$ tyramine feeding experiment, about 4 mg salidroside (**5**) was also isolated. In the feeding experiments using $[1\text{-}^{13}\text{C}]$ phenethylamine, $[1\text{-}^{13}\text{C}]$ 2-phenylethanol, and phenethylamine, a new compound **6** was isolated with HPLC using the same conditions as for the isolation of salidroside. The relative abundance of each ^{13}C -labeled compound was determined by ^{13}C NMR, and the relative incorporation ratio of each deuterium-labeled acteoside was determined by ESI-LC-MS.

2-Phenethyl β -primeveroside (**6**)

Amorphous; negative ESI MS m/z : 415 $[\text{M} - \text{H}]^-$; ^1H NMR δ : 2.92 (2H, dd, $J = 7.5, 7.5$ Hz, H-7), 3.14–3.22 (3H, m, xylose-2, 4, 5a), 3.28–3.36 (3H, m, glucose-2, 3, 4), 3.42 (1H, m, glucose-5), 3.48 (1H, ddd, $J = 10.5, 9.0, 5.5$ Hz, xylose-4), 3.73 (1H, dd, $J = 11.5, 5.5$ Hz, glucose-6a), 3.76 (1H, ddd, $J = 10.0, 7.5, 7.5$ Hz, H-8a), 3.85 (1H, dd, $J = 11.5, 5.5$ Hz, xylose-5b), 4.05 (1H, ddd, $J = 10.0, 7.5, 7.5$ Hz, H-8b), 4.08 (1H, dd, $J = 11.5, 2.0$ Hz, glucose-6b), 4.29 (1H, d, $J = 8.0$ Hz, glucose-1), 4.31 (1H, d, $J = 8.0$ Hz, xylose-1), 7.15 (1H, m, H-4), 7.21–7.28 (4H, m, H-2, 3, 5, 6); ^{13}C NMR δ : 37.2 (t), 66.8 (t), 69.8 (t), 71.1 (d), 71.4 (d), 71.8 (t), 74.8 (d), 75.0 (d), 76.9 (d), 77.6 (d), 77.9 (d), 104.4 (d), 105.4 (d), 127.2 (d), 129.3 (d \times 2), 130.0 (d \times 2), 140.0 (s).

Suppression of acteoside production in olive cells using putative inhibitors

Olive cells were cultured as in the previous experiment. After 2 weeks of additional culturing, 16 mg α -methyltyrosine in 2 ml water (final concentration 1 mM), 24 mg benserazide hydrochloride in 2 ml water (final concentration 1 mM), or 2 ml water was administered into a cell suspension culture. Each compound was sterilized by filtration (Millex-GP 0.22 μm). After administration, each culture was incubated for an additional 2 weeks, after which the cells were harvested and freeze-dried. The dried cells were extracted twice with MeOH at room temperature. The MeOH extracts were concentrated and partitioned between *n*-BuOH and H_2O twice. The *n*-BuOH extracts were concentrated and dissolved up to 10 ml with MeOH. Ten microliters of the solution was injected into the HPLC system fitted with a Shim-pack CLC-ODS(M) (5 μm , C_{18} , 100 \AA , 4.6×250 mm) column and the elution was monitored at 330 nm. The eluate was MeOH/ H_2O /AcOH (50:100:1, v/v/v), the column temperature was 40°C , and the flow rate was 0.5 ml/min. The content of acteoside was

calculated by identifying peak areas. The analysis of each sample was performed in triplicate.

Results and discussion

The induction of olive callus was performed in several modified Murashige and Skoog's agar media. In these media, the callus was induced in CM-NH₄ agar medium supplemented with K⁺ instead of NH₄⁺ and containing 2,4-dichlorophenoxyacetic acid, kinetin, and coconut milk [3]. The cell growth was better in CM-NH₄ medium than in DK-NH₄ medium, which was CM-NH₄ medium without coconut milk. However, the production of acteoside quantified by HPLC analysis was better in a DK-NH₄ liquid medium than in a CM-NH₄ liquid medium (data not shown). Therefore, the induced olive callus was subcultured into the DK-NH₄ agar medium every 4 weeks.

Compounds **1–5** were isolated using HPLC, and identified as acteoside [14], isoacteoside [15], β -oxoacteoside [16], β -hydroxyacteoside [16], and salidroside [7], respectively, using spectra data (Fig. 1). In the olive callus, acteoside was a main secondary metabolite whose content was over 1.7% dry weight (d.w.). This olive cell line will be useful for the provision of acteoside to study biosynthesis or for pharmaceutical activities. The production of phenylethanoid glycosides in olive cell suspension cultures is reported here for the first time, although small amounts of acteoside were isolated from olive fruit [17].

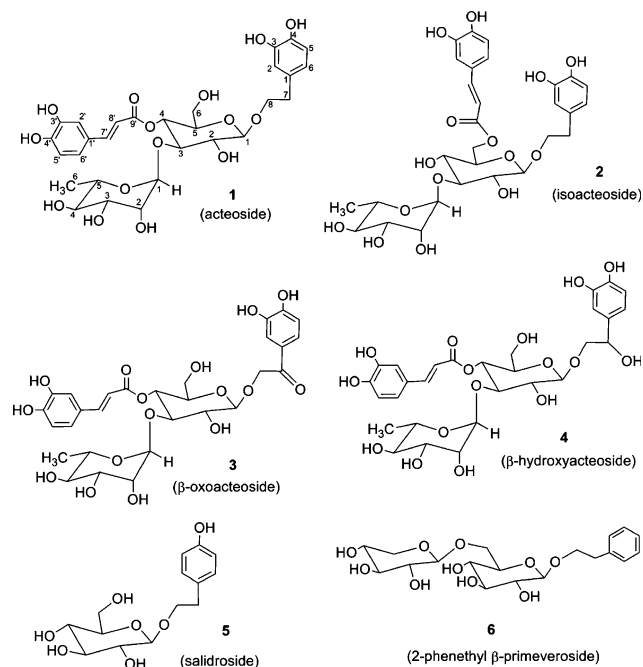


Fig. 1 Structures of compounds **1–6**, isolated from cultured olive cells

Table 2 Relative isotopic abundance of acteoside after ^{13}C -labeled precursor feeding experiments

Carbon position	Precursors			
	A	B	C	D
7	1.1	26.7	12.3	51.3
8	1.1	1.1	12.3	1.1
7'	6.4	1.9	1.1	1.1
8'	1.1	1.1	1.1	1.1
9'	7.3	2.2	1.1	1.1
Glucose-1	1.1	1.1	1.1	1.1

A: phenylalanine ($[1-^{13}\text{C}]:[3-^{13}\text{C}] = 1:1$), 10 mg/flask

B: tyrosine ($[1-^{13}\text{C}]:[3-^{13}\text{C}] = 1:1$), 10 mg/flask

C: $[2,3-^{13}\text{C}_2]$ DOPA, 5 mg/flask

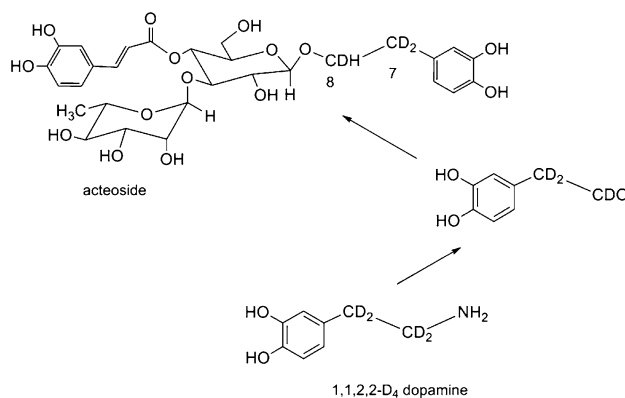
D: $[2-^{13}\text{C}]$ tyramine, 10 mg/flask

We performed labeled precursor feeding experiments to clarify the biosynthetic route to acteoside in olive cells. The ^{13}C -labeled phenylalanine was incorporated into the caffeoyl moiety of acteoside with a 6–7% enrichment at C-7' and C-9' carbons, but not incorporated into the hydroxytyrosol moiety (Table 2). The ^{13}C -labeled tyrosine was incorporated into the hydroxytyrosol moiety with over 25% enrichment at C-7 carbon (Table 2). Therefore, the biosynthetic pathway to the hydroxytyrosol moiety starts from tyrosine, i.e., different from that of the caffeoyl moiety. The labeled tyrosine was also slightly incorporated into the caffeoyl moiety at C-7' and C-9' carbons (Table 2). Phenylalanine ammonia lyase does not generally catalyze tyrosine turnover; however, from our results, phenylalanine ammonia lyase partially catalyzes tyrosine turnover in the olive cells. The $2,3-^{13}\text{C}_2$ -labeled DOPA was incorporated at C-7 and C-8 carbons at about 12% but not into the caffeoyl moiety (Table 2). DOPA would be biosynthesized from tyrosine using tyrosine hydroxylase, as reported in the callus of *Portulaca grandiflora* [18].

The labeled acteosides obtained from feeding experiments using two kinds of deuterium-labeled dopamine were analyzed by LC–ESI–MS. The relative peak intensity and incorporation ratio of each compound are shown in Table 3. $[2,2-^2\text{D}_2]$ Dopamine was incorporated into 20% acteoside with two deuteriums (Table 3). $[1,1,2,2-^2\text{D}_4]$ Dopamine was incorporated into 10% acteoside with three deuteriums or 4% with two deuteriums (Table 3). The results show that the hydroxytyrosol moiety of acteoside is biosynthesized from dopamine. Dopamine was biosynthesized by the decarboxylation of DOPA. $[1,1,2,2-^2\text{D}_4]$ Dopamine was oxidized to 2-(3,4-dihydroxyphenyl)- $[1,2,2-^2\text{D}_3]$ acetaldehyde, reduced to 2-(3,4-dihydroxyphenyl)- $[1,2,2-^2\text{D}_3]$ ethanol, and incorporated into acteoside with three deuteriums. The 2-(3,4-dihydroxyphenyl)- $[1,2,2-^2\text{D}_3]$ ethanol was partially oxidized to 2-(3,4-dihydroxyphenyl)- $[2,2-^2\text{D}_2]$ acetaldehyde and

Table 3 Incorporation of deuterium-labeled dopamines

m/z $[M - \text{H}]^-$	Control	$[2,2-^2\text{D}_2]$ Dopamine	$[1,1,2,2-^2\text{D}_4]$ Dopamine
Relative MS peak intensity of acteoside after feeding deuterium-labeled dopamines			
623	100	100	100
624	29	31	26
625	5.8	31	9.4
626	0	6.8	13
627	0	1.2	3.4
628	0	0	0.6
629	0	0	0
C7–C8			
	Control	$[2,2-^2\text{D}_2]$ Dopamine	$[1,1,2,2-^2\text{D}_4]$ Dopamine
Incorporation ratio of each deuterium			
$\text{CH}_2\text{--CH}_2$	100	80	86
CDH--CH_2	0	0	0
$\text{CD}_2\text{--CH}_2$	0	20	4
$\text{CD}_2\text{--CDH}$	0	0	10
$\text{CD}_2\text{--CD}_2$	0	0	0



reduced to 2-(3,4-dihydroxyphenyl)- $[2,2-^2\text{D}_2]$ ethanol. This is why the acteoside with three or two deuteriums was observed in the feeding experiment of $[1,1,2,2-^2\text{D}_4]$ dopamine.

$[2-^{13}\text{C}]$ Tyramine was enzymatically synthesized from $[3-^{13}\text{C}]$ tyrosine by tyrosine decarboxylase [13]. $[2-^{13}\text{C}]$ Tyramine was incorporated into 50% acteoside at C-7 carbon (Table 2). $[2-^{13}\text{C}]$ Tyramine was also incorporated into salidroside (5), which is a minor phenylethanoid in olive cells (Fig. 2). The production of salidroside was increased 12-fold and labeled almost totally at C-7 carbon when $[2-^{13}\text{C}]$ tyramine was fed to the culture. Therefore, salidroside was biosynthesized from tyramine via *p*-hydroxyphenylacetaldehyde and tyrosol (*p*-hydroxyphenylethanol). It has been reported that, in cell suspension cultures of *Rhodiola sachalinensis*, external tyrosol improved salidroside

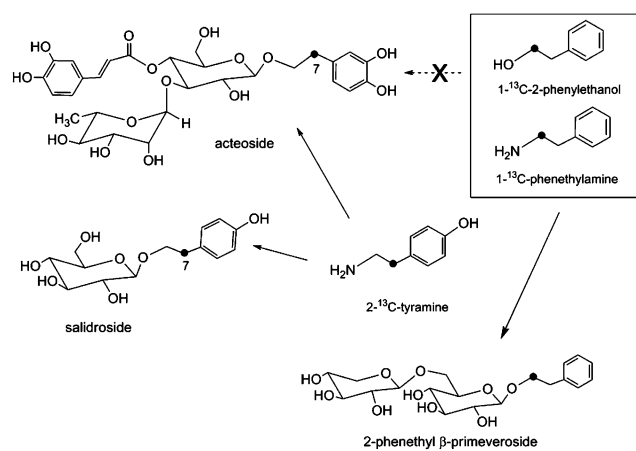


Fig. 2 Incorporation of labeled tyramine, phenylethanol, and phenethylamine in cultured olive cells

production [19]. In cultured olive cells, the ^{13}C -labeled tyramine incorporation into acteoside was also observed, but the incorporation ratio for acteoside was less than that for salidroside. In cell suspension cultures of *Syringa vulgaris*, the incorporation of salidroside to acteoside has been observed [11]. However, based on our results from the tyramine feeding experiment, the incorporation of salidroside into acteoside would be rare in olive cell cultures. The labeled tyramine would be incorporated into acteoside via dopamine or hydroxytyrosol.

In the feeding experiment of benserazide, a DOPA decarboxylase inhibitor, the production of acteoside was reduced to 66%. This result shows that DOPA decarboxylase exists in olives and has an important role in the biosynthesis of acteoside. On the other hand, α -methyltyrosine, an inhibitor of tyrosine hydroxylase, did not reduce acteoside production.

The proposed biosynthetic pathway to acteoside in olive cells is shown in Fig. 3. The hydroxytyrosol moiety of acteoside was biosynthesized from tyrosine through DOPA and dopamine. The amine moiety of dopamine was oxidized to aldehyde, reduced to alcohol, and then β -glycosylated. The olive has alternative biosynthetic pathways available, e.g., from tyramine to acteoside via dopamine or from tyrosol to acteoside via hydroxytyrosol; however, the pathway from tyrosine to acteoside via DOPA and dopamine is the main biosynthetic pathway to acteoside.

[1- ^{13}C]2-Phenylethanol and [1- ^{13}C]phenethylamine were not incorporated into acteoside but incorporated into a new compound **6** which was not detected in the control cell culture. We isolated **6** using HPLC and identified it as 2-phenethyl β -primeveroside (Fig. 1), which is widely detected in various plants [20–22]. The relative isotopic abundance of **6** from [1- ^{13}C]2-phenylethanol and [1- ^{13}C]phenethylamine was over 99%, making **6** the biotransformation product of phenethylamine via phenylethanol in olive cell suspension

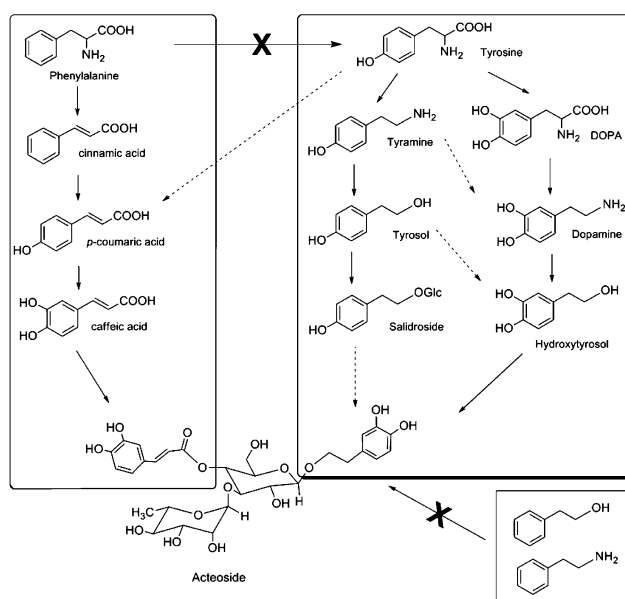


Fig. 3 Proposed biosynthetic pathway to acteoside in cultured olive cells

cultures, not biosynthesized de novo (Fig. 2). From the results of our feeding experiments, the hydroxyl group at C-4 in the benzene ring is important for the biosynthesis of acteoside. Phenylethanoid glycosides are contained in various plant species [23], and the mechanism of glycosylation via the oxidation of amine would be a relatively common reaction in plant secondary metabolism. Further molecular biological research into the glycosylation and total biosynthetic mechanisms of phenylethanoid glycosides is in progress.

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