

(-)-AMPELOPSIN D IS DIFFERENT FROM (-)-QUADRANGULARIN A

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Abstract — The structure of (-)-ampelopsin D, hydroxystilbene dimer, was reinvestigated using spectroscopic methods. Furthermore, the relative and absolute stereostructures were discussed on the basis of chemical transformation of (+)- ϵ -viniferin to (-)-ampelopsin D.

In 1993, Y. Oshima *et al.* reported the isolation and structure of a hydroxystilbene dimer named (-)-ampelopsin D (**1**) as a hepatoprotective substance from the roots of *Ampelopsis brevipedunculata* var. *hancei* (Vitaceae).^{2,3} In 1999, S. A. Adesanya *et al.* reported the isolation and structure of a hydroxystilbene dimer named (-)-quadrangularin A (**2**) from the stems of *Cissus quadrangularis* (Vitaceae).⁴ In their paper, it was described that the NMR data of ampelopsin D (**1**) are quite similar to those of quadrangularin A (**2**) and the reported structure of ampelopsin D is probably erroneous and should be **2**.⁴ Recently, we have isolated a hydroxystilbene dimer (**3**) from the corks of *Vitis vinifera* 'Kyohou' (Vitaceae). The ¹H and ¹³C NMR data of **3** are very similar to those of **1** (the spectra were taken in acetone-d₆) and **2** (the spectra were taken in methanol-d₄). We, therefore, studied the structure of compound (**3**) in detail. In this communication we describe that the relative and absolute structure of (-)-ampelopsin D is **1** and not **2**.

Structure Review of Ampelopsin D

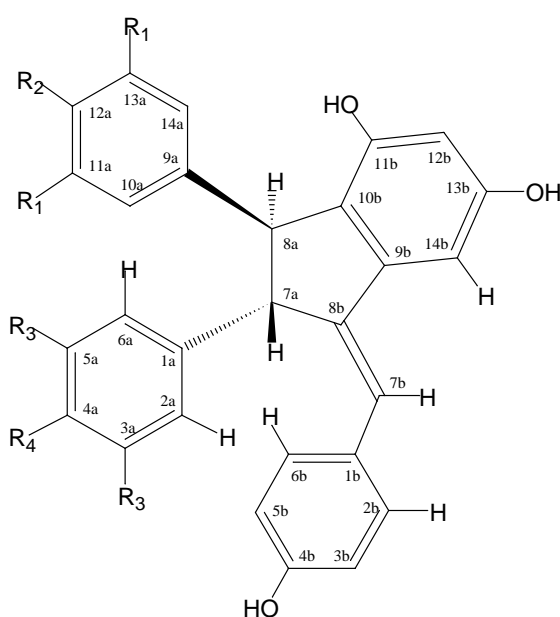
Compound (**3**) isolated from *Vitis vinifera* 'Kyohou', [α]_D²² -5° (c 0.27, MeOH, cell length = 100 mm); HRFABMS *m/z* 455.1498 (calcd for C₂₈H₂₃O₆, 455.1495) showed very similar NMR data to those of (-)-ampelopsin D (**1**) and (-)-quadrangularin A (**2**), respectively.^{1,3} The most important difference between (-)-ampelopsin D (**1**) and (-)-quadrangularin A (**2**) is whether H-2a (H-6a) shows an *ortho*-coupling or a *meta*-coupling. For the structure reviews of **1** and **2**, the structure of compound (**3**) was examined on the

Table 1. NMR Data of Compound (**3**) in Methanol-d₄ (**A**) and in Acetone-d₆ (**B**)

Carbon number	A		B	
	¹³ C	¹ H	¹³ C	¹ H
1a	138.0	s ^a	137.3	s ^a
2a,6a	129.1	d	128.8	d
3a,5a	116.5	d	116.3	d
4a	156.7	s	156.7	s
7a	60.0	d	59.5	d
8a	59.2	d	58.7	d
9a	149.9	s	149.9	s
10a,14a	106.7	d	106.4	d
11a,13a	159.2	s	159.3	s
12a	101.3	d	101.3	d
1b	130.2	s	129.7	s
2b,6b	131.2	d	131.0	d
3b,5b	116.0	d	116.0	d
4b	157.4	s	157.3	s
7b	122.9	d	122.6	d
8b	143.7	s	143.1	s
9b	147.9	s	147.5	s
10b	124.7	s	123.8	s
11b	156.3	s	156.1	s
12b	103.8	d	103.8	d
13b	159.8	s	159.7	s
14b	98.3	d	98.4	d

^amultiplicity of ¹³C signals.

^b*J* (Hz) in parentheses.



(-)-Ampelopsin D (**1**) : R₁=OH, R₂=H, R₃=H, R₄=OH
 (-)-Quadrangularin A (**2**) : R₁=H, R₂=OH, R₃=OH, R₄=H

basis of the NMR spectra taken in methanol- d_4 .⁵ In DIF-NOE experiments of **3**, the NOEs between H-7b and H-14b (4.0%), H-7b and H-2b (H-6b) (2.6%), and H-6b (H-2b) and H-7a (7.0%) were respectively observed. These observations indicated H-7a to be an allylic hydrogen of an Ar - C_{7b} = C_{8b} - C_{7a}-grouping. This was further supported by the following observations. The correlations between H-7a and C-7a in the HMQC spectrum, between H-7b and C-7a in the HMBC spectrum, and between H-7a and H-7b in the ¹H-¹H COSY spectrum, were respectively observed. The long-range coupling between H-7a and H-7b was also confirmed by decoupling experiments.⁶ The broad singlet signal of H-7a changed to a sharp signal by irradiation of H-7b, and the doublet signal of H-7b changed to a singlet signal by irradiation of H-7a. This evidence confirmed the assignments of H-7a and C-7a. Next, the correlations between H-7a and C-2a (C-6a), and between H-2a (H-6a) and C-7a were respectively observed in the HMBC spectrum. The ¹H-¹H coupling constant value of H-2a (H-6a) is 8.8 Hz, which corresponds to a value of *ortho*-coupling.⁷ This means that the structure of compound (**3**) is characterized as **1**.⁸

Absolute Configuration of (-)-Ampelopsin D (+)- ϵ -Viniferin (**4**) ($[\alpha]_D^{22} +49.1^\circ$ (c 1.85, MeOH, cell length = 100 mm), whose absolute configuration is known,⁹ was treated with trifluoromethanesulfonic acid in methanol under reflux for 7 days to give (-)-ampelopsin D (**1**) and its regio-isomer (**5**)¹⁰ based on a double bond, in yields of 7 % and 10 %, respectively. According to the reaction mechanism in Figure 1, (+)- ϵ -viniferin (**4**) will give a product having the structure of **1** and not a product having the structure of **2**, under the above reaction conditions. From the above-mentioned results, (-)-ampelopsin D should be represented as **1**, including the absolute configuration.

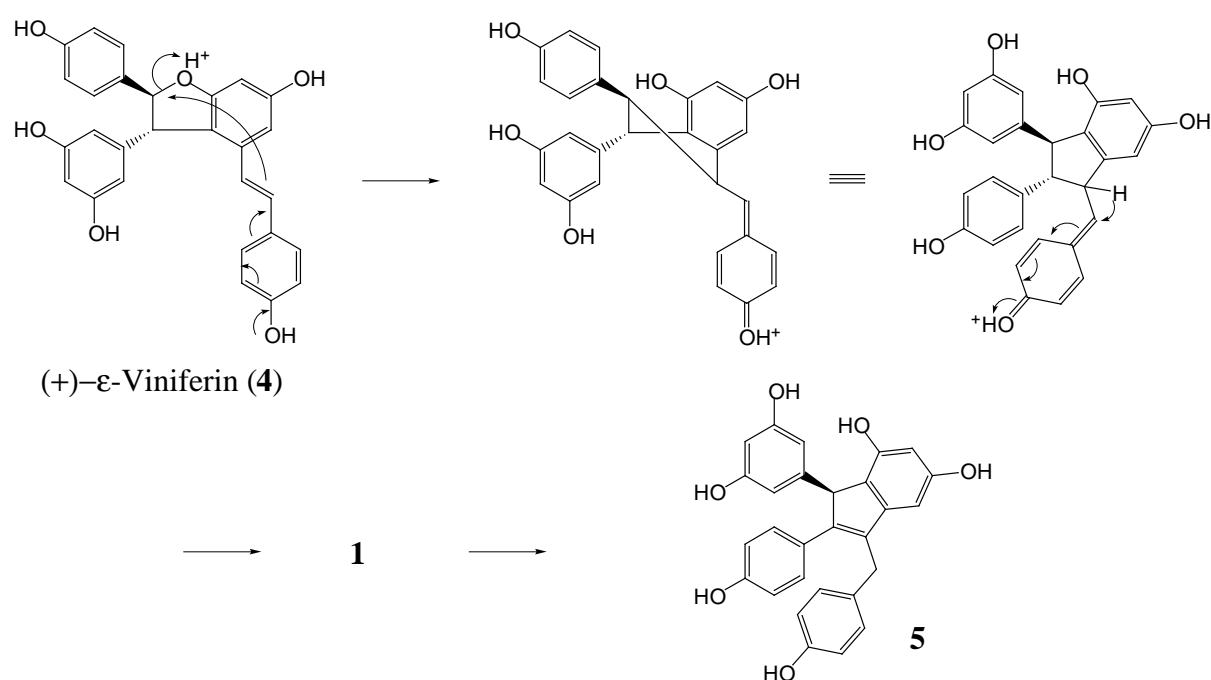


Figure 1. A reaction pathway of (+)- ϵ -viniferin (**4**) to (-)-ampelopsin D (**1**) and its isomer (**5**)

Consequently, the ^1H and ^{13}C NMR data of (–)-ampelopsin D (**1**) and (–)-quadrangularin A (**2**) both are very similar, but (–)-ampelopsin D (**1**) and (–)-quadrangularin A (**2**) are a different compound from each other.

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6. The same long-range coupling was reported by Prof. Y. Oshima *et al.*²
7. The value of H-2a (H-6a) of (–)-quadrangularin A (**2**) was reported as 2 Hz.⁴
8. The ^1H NMR spectrum in acetone- d_6 of **3** was identical with that in acetone- d_6 of **1** sent by Prof. Y. Oshima.
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10. **5**: $[\alpha]_{\text{D}}^{22}$ -227.0° (c 0.39, MeOH, cell length = 100 mm); HRFABMS m/z 455.1499 (calcd for $\text{C}_{28}\text{H}_{23}\text{O}_6$, 455.1495); ^1H NMR (600 MHz, MeOH- d_4) δ 3.81 (1H, d, $J = 16.1$ Hz, H-7b), 3.86 (1H, d, $J = 16.1$ Hz, H-7b'), 4.79 (1H, s, H-8a), 5.98 (1H, t, $J = 2.2$ Hz, H-12a), 6.06 (3H, d, $J = 2.2$ Hz, H-10a, 14a, H-12b), 6.17 (1H, d, $J = 2.2$ Hz, H-14b), 6.65 (2H, d, $J = 8.5$ Hz, H-3a, 5a), 6.72 (2H, d, $J = 8.5$ Hz, H-3b, 5b), 7.06 (2H, d, $J = 8.5$ Hz, H-2a, 6a), 7.10 (2H, d, $J = 8.5$ Hz, H-2b, 6b); ^{13}C NMR (150 MHz, MeOH- d_4) δ 32.2 (t, C-7b), 56.8 (d, C-8a), 100.8 (d, C-14b), 101.1 (d, C-12b), 101.5 (d, C-12a), 108.2 (d, C-10a, 14a), 115.8 (s, C-3a, 5a), 116.3 (d, C-3b, 5b), 125.4 (s, C-10b), 128.9 (s, C-1a), 130.2 (d, 2b, 6b), 131.1 (d, 2a, 6a), 132.1 (s, C-1b), 136.6 (s, C-8b), 144.0 (s, C-9a), 149.9 (s, C-9b), 150.4 (s, C-7a), 154.0 (s, C-11b), 156.5 (s, C-4b), 157.5 (d, C-4a), 158.8 (s, C-11a, 13a), 158.9 (s, C-13b).