



# Erratum

## Corrigendum to “Neolignans from *Piper kadsura* and their anti-neuroinflammatory activity” [Bioorg. Med. Chem. Lett. 20 (2010) 409]

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The authors regret that this article contains a number of errors. The structure of compound **1** was revised. Therefore, the following corrections should be made.

1. In page 409, lines 5–8 from the bottom right of the text should read as: In the positive mode FABMS of **1**, a molecular ion peak [M+H]<sup>+</sup> at *m/z* 357 was observed, and the molecular formula of **1** was determined to be C<sub>20</sub>H<sub>20</sub>O<sub>6</sub> by HR-FABMS and a molecular ion peak [M+H]<sup>+</sup> at *m/z* 357.1335 was observed (calcd for C<sub>20</sub>H<sub>21</sub>O<sub>6</sub>, 357.1338).

2. In page 410, line 5 of the text to the right should read as: one methoxy carbon at  $\delta$  59.2 (OCH<sub>3</sub>-3').

3. In page 411, line 3 of the text to the left should read as: phenyl-7,8-epoxy-8-substituted-propanol group (unit A).

4. In page 411, lines 7–8 of the text to the left should read as: presence of 3-methoxy-4-hydroxy-5-substitued-cyclohexa-1,3-diene group (unit B).

5. In page 411, line 9 of the text to the left should read as: between OCH<sub>3</sub>-3' and C-3' instead of between OCH<sub>3</sub>-4' and C-4'.

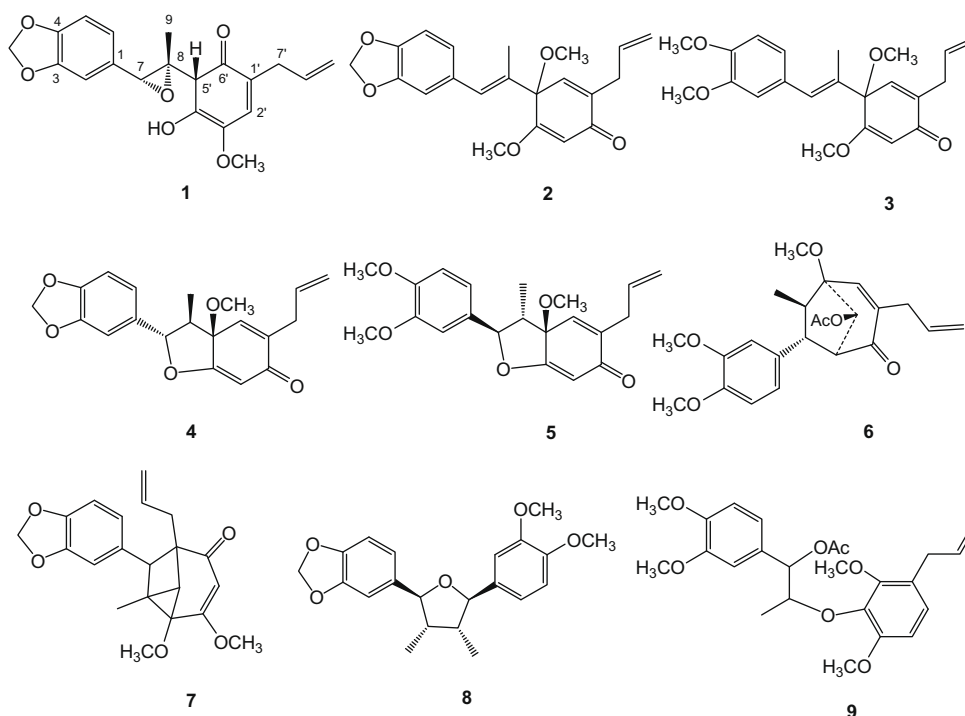


Figure 1. The structures of compounds **1–9** isolated from *P. kadsura*.

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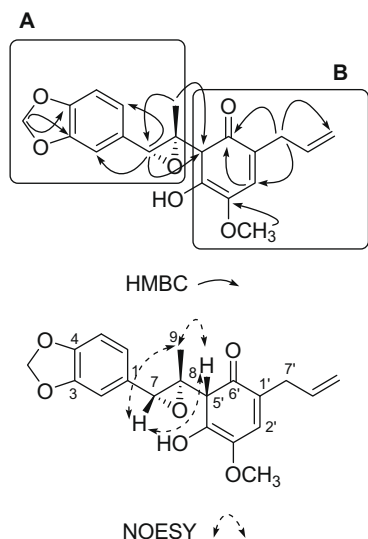
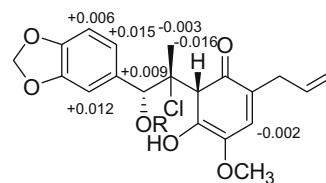
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**Table 1**

<sup>1</sup>H and <sup>13</sup>C NMR data for compound **1** ( $\delta$  in ppm, 500 MHz for <sup>1</sup>H and 125 MHz for <sup>13</sup>C, in CDCl<sub>3</sub>)

Number	<b>1</b>		
	$\delta_H$	$\delta_C$	HMBC
1		133.9, C	
2	6.92 (d, 1.5)	108.8, CH	1, 3, 4, 6, 7
3		147.6, C	
4		146.8, C	
5	6.76 (d, 7.5)	108.1, CH	1, 3, 4, 6
6	6.75 (dd, 1.5, 7.5)	122.7, CH	1, 2, 4, 5, 7
7	3.64 (s)	65.3, CH	1, 2, 6, 8, 9, 4', 5', 6'
8		50.4, C	
9	0.96 (s)	16.8, CH <sub>3</sub>	7, 8, 5', 6'
1'		132.6, C	
2'	7.00 (s)	157.2, CH	1', 3', 4', 6', 7'
3'		131.7, C	
4'		147.0, C	
5'	3.36 (s)	58.6, CH	7, 8, 9, 1', 3', 4', 6'
6'		197.2, C	
7'	2.91 (dd, 1.5, 5.5)	33.0, CH <sub>2</sub>	1', 2', 6', 8', 9'
8'	5.82 (m)	135.5, CH	1', 7', 9'
9'	5.06 (dd, 1.5, 5.5)	116.7, CH <sub>2</sub>	1', 7', 8'
	5.08 (dd, 1.5, 12.0)		
O-CH <sub>2</sub> -O	5.94 (s)	101.1, CH <sub>2</sub>	3, 4
OCH <sub>3</sub> -3'	3.82 (s)	59.2, CH <sub>3</sub>	3'

Assignments were confirmed by <sup>1</sup>H-<sup>1</sup>H COSY, HMQC, HMBC, and NOESY spectra.

**Figure 2.** Key 2D NMR (HMBC, NOESY) correlations of **1**.

**1s** R = (S)-MTPA

**1r** R = (R)-MTPA

**Figure 3.** Difference in the  $\Delta_{SR}$  ( $\delta_S - \delta_R$ ) values of the chlorinated MTPA esters of **1**.

6. In page 411, lines 11–13 of the text to the left should read as: in the HMBC spectrum from H-7 to C-4', C-5', C-6', from H-9 to C-5', C-6', and from H-5' to C-7, C-8, C-9.

7. In page 411, lines 3–5 of the text to the right should read as: Compound **1** was fortunately, esterified by (S)- and (R)-MTPA chlorides to yield the chlorinated (R)- and (S)-MTPA esters, respectively.

8. In page 411, lines 8–9 of the text to the right should read as: C-7 of **1** was the R configuration. We therefore propose the structure of **1** in Figure 1, which is similar to that of piperkadsin B.<sup>2</sup>

9. In page 412, lines 1–5 of the text to the left should read as: compound **1** represents a rearranged neolignan of piperkadsin B bearing 7,8-epoxy group. Thus, compound **1** was determined to be a new neolignan, (7R)-7,8-epoxy-3,4-methylenedioxyphenyl-3'-methoxy-4'-hydroxy-6'-oxo- $\Delta$ -1',3',8'-8,5'-lignan, namely, piperkadsin C.

10. In Supplementary Content:

**Piperkadsin C (1):** FAB-MS (positive mode)  $m/z$ : 357 [M+H]<sup>+</sup>; HRFABMS  $m/z$ : 357.1335 [M+H]<sup>+</sup> (Calcd for C<sub>20</sub>H<sub>21</sub>O<sub>6</sub>: 357.1338).

11. There were errors in Table 1. The correct format of Table 1 appears below.

12. In Figure 1 was corrected. The correct format of Figure 1 appears below.

13. In Figure 2 was corrected. The correct format of Figure 2 appears below.

14. In Figure 3 was corrected. The correct format of Figure 3 appears below.