



Immunosuppressive chalcone-isoflavonoid dimers from *Campylotropis hirtella*

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ABSTRACT

Twelve new chalcone-isoflavonoid dimers were isolated from the roots of *Campylotropis hirtella*. The structures of these compounds were elucidated by NMR analysis, while the absolute configurations were determined from the CD spectra. We measured immunosuppressive activities of these compounds on splenic lymphocytes and found that many of the compounds displayed immunosuppressive activities. Compound **8** was the most active, with IC₅₀ values of 3.68 μM for T lymphocyte suppression and 1.79 μM for B lymphocyte suppression.

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1. Introduction

Flavonoids are a class of plant metabolites often studied for their pharmacological activities, such as antiviral, anti-inflammatory and antitumor effects. These natural polyphenolic compounds, with a carbon skeleton of C6–C3–C6, can easily undergo phenol oxidative coupling reactions and aggregate into bioflavonoids (C6–C3–C6)₂. Due to diversity of flavonoids, its coupling product, biflavonoids can also be classified into various classes.¹ Within biflavonoid classes, the natural occurring arylbenzofuran-isoflavonoid dimers was rarely reported; only few papers described its presence in *Lespedeza cyrtobotrya* Miq. and *Lespedeza floribunda*.² Our group has previously reported that many novel isoflavonoids separated from the roots of *Radix Campylotropis hirtellae* show strong immunosuppressive activity.³ However, biflavonoids from this plant have never been studied. To further explore naturally occurring immunosuppressive agents, 12 new chalcone-isoflavonoid dimers were isolated from *C. hirtellae*. The structures of these compounds were elucidated by NMR analysis, while the absolute configurations were determined from the CD spectra. The carbonyl bridge between 2-arylbenzofuran and isoflavonoid subunit, crucial in initiating phenol oxidative coupling reaction, plays

a key role in determining interflavonoid bond location. By measuring immunosuppressive activities on splenic lymphocytes, we found that many compounds were immunosuppressive and that compound **8** was the most active, with IC₅₀ values of 3.68 μM for T lymphocyte suppression and 1.79 μM for B lymphocyte suppression.

2. Results and discussion

Hirtellane A–E (**1–5**) were assumed to be dimeric flavonoids composed of an isoflavan and a 2-arylbenzofuran from their molecular formula and NMR data. The HR-ESI-MS of compound **1** (Hirtellane A) displayed a molecular ion peak at 651.1878, suggesting a molecular formula of C₃₇H₃₂O₁₁. The ¹³C NMR spectra showed a total of 37 signals for **1** (Table 1). The ¹H NMR spectrum and HMQC experiments showed 29 sp² carbons, including 11 carbons bearing an oxygen atom and one carbonyl, and eight sp³ carbons corresponding to two aromatic methoxyl groups, two geminated methyl groups, two methylene, one methine and a quaternary carbon. The ¹H NMR spectrum of Hirtellane A showed five aliphatic proton signals at δ_H 2.51 (1H, d, 15.6 Hz), δ_H 2.73 (1H, dd, 15.6 Hz, 12.4 Hz), δ_H 3.12 (1H, brs), δ_H 3.69 (1H, d, 10.2 Hz) and δ_H 4.03 (1H, d, 10.2 Hz), which were characteristic isoflavan signals.⁴ The ¹H NMR also showed a set of 2, 2-dimethylpyrene ring protons at δ_H 1.35, δ_H 1.38 (each 3H, s), δ_H 6.48 and δ_H 5.58 (each 1H, 10 Hz). In addition, the olefinic signal at δ_H 6.48 coupled with methoxyl signal at δ_H 3.65 in NOE

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Table 1¹H and ¹³C NMR Spectroscopic data of compound **1**–**4** in Acetone-*d*₆

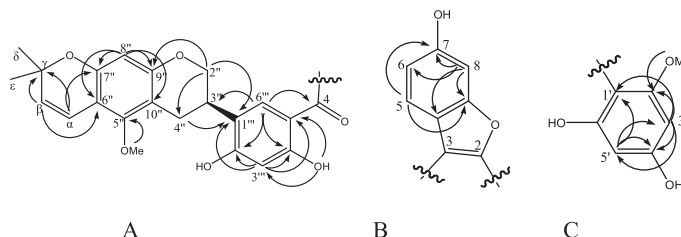
Position	Compound 1	Compound 2	Compound 3	Compound 4
	δ_{H} (J in Hz)	δ_{C}	δ_{H} (J in Hz)	δ_{C}
2		151.8	150.9	150.8
3		117.2	117.0	116.9
4		195.1	194.5	194.2
5	7.58 (1H, d, 8.5 Hz)	121.3	7.20 (1H, s)	104.7
6	6.90 (1H, dd, 8.5 Hz, 1.9 Hz)	112.7	145.1	142.4
7		156.0	145.3	144.1
8	6.99 (1H, d, 1.7 Hz)	97.7	7.08 (1H, s)	97.1
9		155.6	148.8	148.4
10		120.5	118.8	119.3
1'		99.5	98.9	103.4
2'		159.6	157.1	157.9
3'	6.14 (1H, s)	96.1	6.14 (1H, d, 2.0 Hz)	113.4
4'		162.1	160.4	158.0
5'	5.91 (1H, s)	91.4	5.93 (1H, d, 2.0 Hz)	98.7
6'		157.9	158.8	154.5
2''	3.69 (1H, d, 10.2 Hz)	69.4	3.74 (1H, m)	68.7
	4.03 (1H, d, 10.3 Hz)		4.04 (1H, m)	
3''	3.12 (1H, brs)	32.1	3.15 (1H, brs)	31.2
4''	2.51 (1H, t, 15.6 Hz)	25.1	2.54 (1H, t, 15.6 Hz)	24.0
	2.73 (1H, dd, 15.6 Hz, 12.4 Hz)		2.74 (1H, m)	
5''		152.9	152.3	152.2
6''		107.9	107.2	107.2
7''		154.9	154.2	154.2
8''	5.99 (1H, s)	100.1	6.00 (1H, s)	99.4
9''		155.7	155.0	155.1
10''		108.7	102.0	107.9
1'''		119.2	118.5	118.3
2'''		162.1	161.6	161.6
3'''	6.44 (1H, s)	102.6	6.48 (1H, s)	102.1
4'''		164.1	163.3	163.8
5'''		113.8	113.1	113.0
6'''	7.44 (1H, s)	132.2	7.46 (1H, s)	131.5
α	6.48 (1H, d, 10.0 Hz)	117.2	6.51 (1H, d, 10.0 Hz)	116.5
β	5.58 (1H, d, 10.0 Hz)	127.8	5.59 (1H, d, 10.0 Hz)	127.1
γ		75.8	75.1	75.0
δ	1.38 (3H, s)	27.5	1.40 (3H, s)	26.7
ϵ	1.35 (3H, s)	27.1	1.40 (3H, s)	26.7
α'			3.11 (1H, t, 5.8 Hz)	21.8
β'			4.96 (1H, t, 6.8 Hz)	123.2
γ'				129.2
δ'			1.65 (3H, s)	24.5
ϵ'			1.63 (3H, s)	16.5
2'-OMe	3.52 (3H, s)	61.1	3.51 (3H, s)	60.3
5'-OMe	3.65 (3H, s)	54.9	3.68 (3H, s)	54.1
5-OMe		3.94 (3H, s)		55.5
OH	12.66		12.82	
	8.80,	8.85	8.6–8.9	
	8.72			12.64
				9.93,
				9.14

experiment.^{1a} In HMBC, a chelated hydroxyl signal at δ_{H} 12.8 coupled with C-4''' (δ_{C} 164.1), C-5''' (δ_{C} 113.8) and C-3''' (δ_{C} 102.6). A singlet signal at δ_{H} 7.44 (δ_{C} 132.2) was long-range coupled with C-4 (δ_{C} 195.1), C-4''' (δ_{C} 164.1), C-2''' (δ_{C} 162.1) and C-3'' (δ_{C} 32.1). The singlet aromatic proton signal at δ_{H} 6.44 (δ_{C} 102.6) coupled with C-4''' (δ_{C} 164.1), C-2''' (δ_{C} 162.1), C-1''' (δ_{C} 119.2) and C-5''' (δ_{C} 113.8). These data led to the assignment of the substructure A (Fig. 1).

In ¹H NMR, three aromatic protons δ_{H} 7.58 (1H, d, 8.5 Hz), δ_{H} 6.90 (1H, dd, 8.5 Hz, 1.9 Hz) and δ_{H} 6.99 (1H, d, 1.9 Hz) form a typical ABX pattern. In HSQC, δ_{H} 6.99 corresponding to δ_{C} 97.7 and in HMBC, this proton was long-range coupled to C-9 (δ_{C} 155.3), C-7 (δ_{C} 156.3), C-6 (δ_{C} 112.7) and C-10 (δ_{C} 120.5). The aromatic proton signal at δ_{H} 6.14 was coupled to C-4' (δ_{C} 161.1), C-1' (δ_{C} 99.5) and C-3' (δ_{C} 91.4). The signal at δ_{H} 5.91 was coupled to C-4' (δ_{C} 161.1), C-2' (δ_{C} 159.6), C-1' (δ_{C} 99.5) and C-5' (δ_{C} 96.1). A methoxyl signal at δ_{H} 3.52 correlated to C-2' (δ_{C} 159.6). These data led to establish the substructure B and C (Fig. 1).⁵ Thus, the two reminiscent tetra-substituted sp² carbons (δ_{C} 151.8 and δ_{C} 117.5) should be involved in

the linkage of these three substructures. Therefore, four alternative arylbenzofuran structures (**1a**–**1d**, Fig. 2) could be proposed for **1**.

Only five chalcone-flavone dimers⁶ and eight pterocarpan-benzofuran dimers² with the carbon skeletons **1a** and **1c** have already been described in the literature, and the distinct chemical

**Fig. 1.** Key HMBC (H→C) correlations of compound **1**.

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