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OPEN Glaulactams A–C, daphniphyllum alkaloids from Daphniphyllum glaucescens

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Glaulactams A-C (1-3), which possess a novel skeleton, as well as the known compound daphmanidin B (4), were isolated from the leaves of Daphniphyllum glaucescens and separated using ion-exchange chromatography aided by NMR fingerprinting. Their structures, including their absolute configurations, were elucidated by spectroscopic analyses and time-dependent density-functional-theory-calculated electronic circular dichroism spectra; the data were subsequently analyzed to gain insight into the respective biogenetic relationships between the isolates, which exhibited anti-H1N1 and immunosuppressive activities.

The first daphniphylline-type alkaloid, daphniphylline, which possesses a polycyclic, C₃₀-aliphatic structure, was discovered in 1966¹. A number of novel structures from the Daphniphyllaceae family of plants were recently reported, including himalensine A, with its 13,14,22-trinorcalyciphylline A backbone², himalensine B with its 22-nor-1,13-secodaphnicyclidin framework², and macropodumines A-C with fused pentacyclic ring systems³. The complex and fascinating structures of these alkaloids render them synthetically challenging⁴. Many types of alkaloid are reported to be produced by Daphniphyllum glaucescens, including daphniglaucin C, which has a novel structure that contains hexahydroazulene and octahydroindole ring systems⁵ and daphniglaucins A and B, which have unique 1-azoniatetracyclo[5.2.2.0^{1,6}0.^{4,9}] undecane motifs⁶. In addition, daphniglaucins D-H possess fused hexacyclic skeletons, and daphniglaucins J and K are yuzurimine-type alkaloids that have previously been reported from this species⁷.

Natural product chemists are very interested in the discovery of novel structures with unique properties⁸. To date, a variety of methods have been used to facilitate the discovery of novel natural products, with examples including NMR-fingerprinting⁸ and LC-MS-guided methods⁹. Daphnezomine-F-type^{10,11} and daphmanidin-C-type¹² alkaloids are rare groups of daphniphyllum alkaloids; they each possess a lactam functional group that is biogenetically derived through the oxidative cleavage of a yuzurimine C-C bond. The presence of lactams in these daphniphyllum compounds inspired us to develop a new method that combines ion-exchange chromatography (IXC) and NMR fingerprinting to screen for new alkaloids. Hence, in this article, we report the isolation, structural characterization, and biological evaluation of glaulactams A-C (1-3) from the leaves of Daphniphyllum glaucescens.

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Figure 1. Structures of compounds 1-3.

Results and Discussion

Air-dried *D. glaucescens* leaves (1.53 kg) were extracted with MeOH and concentrated to obtain a crude extract, which was dispersed in 80% aqueous MeOH to give a methanolic suspension. This suspension was then partitioned with hexane $(4 \times 2 \text{ L})$ to remove the lipid constituents. The methanolic extract was filtered and passed through a cation-exchange resin. The eluent containing the acidic and neutral compounds was concentrated under reduced pressure and subjected to Diaion HP20 column chromatography, followed by vacuum liquid chromatography (VLC) over silica gel to yield 26 fractions (4A-4Z); among these fractions, the ¹H NMR spectra of fractions 4V, 4W, and 4Z were found to exhibit signals characteristic of daphniphyllum alkaloids (see the Extraction and Isolation section). As a result, these three fractions were selected for repeated column chromatography over silica gel and RP-18 stationary phases to yield three compounds that had structures that were consistent with 1 (4.2 mg), 2 (29.2 mg), and 3 (3.0 mg) (Fig. 1), as described below.

Glaulactam Å (1) exhibited a molecular ion peak at m/z 486.2494 [M + H]⁺ (calcd., 486.2486), consistent with the $C_{27}H_{35}NO_7$ molecular formula and 11 degrees of unsaturation. The IR spectrum of 1 exhibited bands corresponding to carbonyl functionalities (1736, 1730, 1712, and 1695 cm⁻¹). The ¹³C and ¹H NMR data (Table 1) reveal the presence of 27 carbon signals ascribed to two acetyl groups (δ_C 168.9, 20.1, δ_H 2.05; δ_C 170.5, 20.4, δ_H 2.07), a methyl carboxylate (δ_C 173.9, 50.4, δ_H 3.59), a conjugated diene (δ_C 120.9, 134.1, 134.6, 143.3, δ_H 5.35), an acetoxy-bearing methine (δ_C 166.4, δ_H 4.06, 5.03), and an amide carbonyl (δ_C 180.5). The above functionalities accounted for six out of the 11 degrees of unsaturation, suggesting that 1 is pentacyclic. The HSQC, ¹H–¹H COSY, and HSQC-COSY spectra of 1 reveal three structural connectivities, namely I (H-2 to H-18, H-2 to H-4, and H-18 to both H₂-19 and H₃-20), II (H-6 to H₂-7, H-11 to H₂-12, and H-6 to H₂-12), and III (H₂-13 to H₂-17), as is illustrated in Fig. 2.

Subsequent HMBC spectral analysis revealed where these three structural fragments are linked. The C-1 carbonyl ($\delta_{\rm C}$ 180.5) and C-19 ($\delta_{\rm C}$ 51.4, $\delta_{\rm H}$ 3.16, m, 2 H) of fragment I were speculated to be due to a γ -lactam ring, which was verified by the observed HMBC-derived H₂-3/C-1 and H₂-19/C-1 correlations. HMBC correlations between H₂-21 and C-4, C-5, C-6, and C-8, and between H₂-7 and C-1 and C-19 linked fragment I to II, resulting in an amide-bridged bicyclic system (C-1 to C-7 and the N atom, and C-18 to C-19). The link between the methyl carboxylate and C-14 in fragment III was made on the basis of the HMBC correlations between both H-14 and H₃-23, and C-22, while C-13, C-15, and C-17 are attached to C-8, C-9, and C-10 of the conjugated diene on the basis of the HMBC correlations between H-14 and C-8 and C-9, and between H-11 and C-9 and C-17. The acetoxy groups at C-4 and C-21 were assigned on the basis of HMBC correlations between H-4 and H₂-21 and the respective acetyl carbonyls at $\delta_{\rm C}$ 168.9 and 170.5. Hence, the planar structure of 1 was assigned to be a fused pentacyclic ring with a γ -lactam functionality (Fig. 2).

The relative configuration of 1 was deduced through NOE-correlation analysis, as indicated in the Chem3D drawing (Fig. 3); H-4/H-19, H-4/H-6, H-4/H₃-20, H-21a/H-12b, and H-6/H₂-12 NOE correlations indicate that H₂-21, H-6, H₃-20, and H-4 are cofacial and were arbitrarily assigned to be β -oriented. In addition, the NOE spectrum reveals a correlation between H-2 and H₂-3, in addition to a correlation between H-3a and H₃-20, implying that H-2 is positioned between H₂-3, but opposite to H₃-20 (Fig. 3). In addition, H-13a/H-21b, H-13b/H-14, and H-14/H-15 correlations suggest that both H-14 and H-15 are α -oriented.

High-resolution electrospray-ionization mass spectrometry (HRESIMS) revealed that glaulactam B (2) has the molecular formula $C_{25}H_{33}NO_5$, the mass of which is less than that of 1 by one CH_2CO_2 unit. In addition, the ¹³C and ¹H NMR data for 2 are similar to those of 1; however, instead of possessing an acetoxy-bearing methine, as in 1, an sp³ methylene is present at C-4 (δ_C 36.3) in 2 (Table 1). This was confirmed by the HMBC correlations between H_2 -21 and C-4, C-5, C-6, and C-8 (Fig. 2). The planar structure of 2 (Fig. 2) was established following ¹H-¹H COSY, HSQC-TOCSY, and HMBC spectral analyses. With the exception of C-4, the same relative configurations were assigned to 1 and 2 on the basis of detailed NOE-correlation analyses (Supplementary Fig. S4).

		1		2		3	
no.		¹³ C	¹ H (J in Hz)	¹³ C	¹ H (J in Hz)	¹³ C	¹ H (J in Hz)
1		180.5 s		181.6 s		181.7 s	
2		46.1 d	2.37 m	45.8 d	2.27 m	45.5 d	2.15 m
3	a	35.9 t	2.03 m	28.9 t	1.92 m	27.7 t	1.87 m
3	b		2.75 m		2.55 m		2.91 m
4	a	76.6 d	5.51 dd (11.7, 3.6)	36.3 t	1.93 m	35.2 t	2.19 m
	b				2.16 m		2.27 m
5		51.0 s		45.9 s		45.2 s	
6		40.3 d	3.12 m	41.8 d	2.85 m	41.3 d	2.97 m
7	a	45.8 t	2.59 dd (13.3, 3.2)	46.0 t	2.56 dd (13.1, 3.3)	43.9 t	3.83 dd (13.3, 3.7)
	b		4.65 t (13.3)		4.66 t (13.1)		4.67 t (13.3)
8		134.1 s		136.4 s		126.7 s	
9		143.3 s		142.1 s		148.7 s	
10		134.6 s		134.6 s		168.2 s	
11		120.9 d	5.35 m	120.3 d	5.32 br d (5.4)	27.7 t	2.37 m
							2.61 m
12	a	33.2 t	2.06 m	32.9 t	1.99 m	25.9 t	1.68 m
	b		2.72 m		2.63 m		2.22 m
13	a	42.4 t	3.19 d (14.5)	42.8 t	2.90 d (15.2)	134.2 d	7.03 s
	b		3.76 m		3.72 m		
14		47.2 d	3.27 t (7.9)	46.6 d	3.21 dd (7.6, 7.3)	119.4 s	
15		54.3 d	3.69 m	54.3 d	3.67 m	160.2 s	
16	a	27.7 t	1.09 m	27.8 t	1.11 m	43.6 t	2.74 m, 2H
	b		1.74 m		1.73 m		
17	a	38.5 t	2.41 m	38.4 t	2.41 dd (14.4, 6.8)	25.5 t	2.77 m
	b		2.57 m		2.59 m		2.84 m
18		28.1 d	2.33 m	28.0 d	2.30 m	27.9 d	2.29 m
19	a	51.4 t	3.16 m, 2H	51.8 t	3.13 t (9.5)	51.6 t	3.17 t (9.7)
	b				3.16 t (9.5)		3.22 t (9.7)
20		12.5 q	1.25 d (6.7)	12.6 q	0.97 d (6.7)	12.4 q	1.00 d (7.0)
21	a	66.4 t	4.06 d (11.6)	73.0 t	4.13 d (11.0)	77.0 t	3.91 d (11.0)
	b		5.03 d (11.6)		4.40 d (11.0)		4.47 d (11.0)
22		173.9 s		173.9 s		164.9 s	
23		50.4 q	3.59 s	50.3 q	3.59 s	50.3 q	3.75 s
4-OAc		168.9 s					
		20.1 q	2.05 s				
21-OAc		170.5 s		170.8 s		170.2 s	
		20.4 q	2.07 s	20.4 q	2.15 s	20.3 q	2.13 s

Table 1. ¹³C and ¹H NMR spectroscopic data for 1–3 in pyridine- d_5 (125/500 MHz).

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Glaulactam C (3) was subjected to HRESIMS, which suggested that its molecular formula was $C_{25}H_{31}NO_5$, which contains two less hydrogen atoms than 2. IR absorptions at 1735, 1701, 1687, and 1641 cm⁻¹ indicate the presence of carbonyl and alkene functionalities. The ¹H and ¹³C NMR spectra of 3 reveal the presence of characteristic signals that are attributable to a similar γ -lactam ring (δ_C 181.7, 45.5, 27.9, 51.6, and 12.4), a 21-acetoxy moiety (δ_C 77.0, 170.2, and 20.3), and a 14-methyl carboxylate (δ_C 164.9, 50.3) functionality as was observed for 2. Moreover, the presence of six olefinic-carbon signals (δ_C 119.4, 126.7, 134.2, 148.7, 160.2, and 168.2; δ_H 7.03) and an upfield-shifted carboxylate carbon (δ_C 164.9) that is correlated to the methyl protons (δ_H 3.75) in the HMBC spectrum, suggest that the carboxylate is conjugated to the three alkene units (Fig. 2). The HMBC correlations between H-13 and C-8, C-9, C-14, and C-15, and between H₂-16 and C-9 and C-10, as well as H₂-11 and C-9, C-10, and C-17, led to the assignment of a fulvene moiety, as illustrated in Fig. 2. The planar structure of 3 was confirmed by further HSQC-TOCSY spin-system analyses (H-6 to H₂-7, H-11 to H₂-12, and H-6 to H₂-12) and HMBC correlations between H₂-21 and C-4, C-5, C-6, and C-8. The H-21b/H-6, H-19b/H₃-20, and H-19b/H-6 NOE correlations imply that H-6, H₂-21, and H₃-20 are oriented on the same face of the molecule. Moreover, H₃-20/H-3a, H-2/H-3b correlations suggest that H-2 is positioned opposite to H₃-20 (Fig. 3).

The absolute configurations of compounds 1–3 were determined by comparing the experimental electronic circular dichroism (ECD) spectra to those calculated theoretically. Compounds 1–3 were subjected to standard conformational analysis as implemented in the Confab program¹³. The generated lowest-energy structures following further B3LYP/6–31G(d) optimizations were then used to calculate ECD spectra by time-dependent density functional theory (TDDFT) at the TD-CAM-B3LYP/def2TZVP level. All calculations were performed





using the Gaussian 09 Rev. D program package using the "ultrafine grid" option, *Integral (Grid = UltraFine)*, with solvent effects accounted for using the IEFPCM method¹⁴. The ECD spectrum of each compound 1–3 was finally generated as the Boltzmann-weighted sum of the spectra generated for the various conformers in each case, which resulted in the establishment of the absolute configurations of glaulactams A–C were, as illustrated by structures 1–3, since the calculated spectra are in good agreement with those acquired experimentally (Fig. 4; Supplementary Fig. S5).

It is possible to describe the biosynthetic origins of 1–4 on the basis of the lactam formation mechanism of hemiaminals, the structure of yuzurimine E^{15} and a proposed compound i. In the proposed mechanism (Fig. 5), oxidative cleavage of the C-1–C-2 bond in yuzurimine E via pathway a^{16} yields intermediate ii, which, after oxidation at C-2, gives 4. Similarly, C-1–C-8 bond cleavage via pathway *b* results in the formation of intermediate iii; subsequent deprotonation of the iminol affords lactam 2, which then undergoes oxidation at C-4 and acetylation to yield 1. Double-bond migration and dehydrogenation then result in the transformation of 2 into fulvene 3.

In preliminary biological screening, the extract of D. glaucescens exhibited antiviral activity against the influenza virus and was immunosuppressive in lipopolysaccharide (LPS)-stimulated murine dendritic cells (DCs). Compounds 1-4 were tested for their anti-influenza virus (i.e., anti-H1N1) activities in Madin-Darby canine kidney (MDCK) cells using the plaque assay with betulinic acid as the positive control¹⁷. Cytotoxicity testing revealed that the isolates were not toxic to uninfected host MDCK cells at a concentration of $100 \,\mu M$ (Fig. 6a). However, at a concentration of 50 µM, compounds 1 and 4 were found to substantially inhibit plaque formation of MDCK cells by H1N1 virus infection, to values of 24.4% and 28.0%, respectively. Although less effective under the same treatment conditions, compounds 2 and 3 were found to moderately inhibit plaque formation (69.1% and 63.5%, respectively) (Fig. 6b). In addition, compounds 1-3 were also evaluated for their immunosuppressive activities. Mouse bone-marrow DCs were treated with compounds 1-3, and the immunosuppressive agent quercetin was used as the positive control¹⁸. In preliminary studies, compounds 1-3 (50 µg/mL) and quercetin (50 μ M) were found to have no significant cytotoxic effects on murine DCs in the presence of LPS (100 ng/ mL) (Fig. 7). However, subsequent experiments revealed that compounds 1–3 significantly suppress the levels of tumor necrosis factor- α (TNF- α), interleukin-6 (IL-6), IL-12p70, and nitric oxide (NO) in LPS-stimulated murine DCs (Fig. 7). These results confirm that the immunosuppressive properties of compounds 1–3 are not due to their cytotoxicities in DCs, and that the observed effects are similar to that induced by quercetin.

In summary, IXC and NMR fingerprinting were used to identify and isolate three novel daphniphyllum alkaloids, whose anti-influenza and immunosuppressive activities were then explored. The method described herein can be implemented as a convenient alternative to existing methods commonly employed to extract unique compounds from a complex array of natural products.

Methods

Ethical Statement. The Institutional Animal Care and Use Committee (IACUC) of National Chung Hsing University approved the experimental procedures (approved protocol no. NCHU-IACUC-104-027). The methods were performed in accordance with the approved guidelines.

General experimental procedures. Optical rotations were measured on a JASCO P2000 digital polarimeter and IR spectra were acquired on a Shimadzu IR Prestige-21 FT-IR spectrometer. NMR spectra, in pyridine- d_5 , were recorded on a 500 MHz Avance III spectrometer (Bruker, Rheinstetten, Germany). The ¹H and ¹³C NMR chemical shifts were referenced to the solvent residual peaks at δ_H 7.58 and δ_C 135.5 for pyridine- d_5 . HRESIMS was performed on an LTQ Orbitrap XL mass spectrometer. Silica gel 60 (Merck, 230–400 mesh), Diaion HP-20 (SupelcoTM, Bellefonte, PA, USA), and RP-18 gel (LiChroprep 40–63 µm, Merck) were used for column chromatography. A Shimadzu LC-20AT pump and Shimadzu SPD-M20A diode array detector (Shimadzu Inc., Kyoto, Japan) equipped with an Inertsil ODS-3 column (5 µm, 250 × 10 mm, GL Science Inc., Tokyo, Japan) were



Figure 3. Selected NOESY correlations in 1 and 3.







Figure 5. Plausible biogenetic pathways for 1-4.



Figure 6. Cytotoxic (**a**) and anti-H1N1(**b**) activities of compounds 1–4. Betulinic acid was used as the positive control.

used for HPLC. Preparative medium-pressure liquid chromatography (MPLC) was performed on an Interchim PuriFlash XS 420 chromatographic system.

Plant Material. The leaves of *D. glaucescens* subsp. *oldhamii* were collected in Yilan County, Taiwan in October 2014. The plant was identified by Dr. S.-Y. Hwang. A voucher specimen (specimen no. DG-Chao004) was deposited at the Chinese Medicine Research and Development Center, China Medical University Hospital.

Extraction and Isolation. Air-dried leaves of *D. glaucescens* (10.38 kg) were exhaustively minced and extracted with MeOH (4×20 L). The solvent was concentrated, and 20% water was added to yield an aqueous suspension, to which hexane (4×2 L) was added to removed lipids and chlorophylls. The methanolic solution





was passed through a strong cation-exchange resin (Dowex 50WX4) to adsorb alkaloids (Supplementary Fig. S6). The collected eluent, which contained nonalkaloid constituents (1530 g), was fractionated by column chromatography (CC; Diaion HP-20: column size: 12.0×25.0 cm; successively eluted with 10 L of 30%, 60%, 80%, and 100% MeOH in H_2O to yield four fractions. Fraction 4 (24.95g), which was eluted with 100% MeOH, was fractionated by silica-gel CC (column size: 6.0×15.0 cm; eluted with 9:1 to 0:1 hexane:EtOAc) to yield 26 subfractions (4A-4W). Subfraction 4V (203 mg), which exhibited ¹H NMR signals characteristic of daphniphyllum alkaloids (i.e., few methyl signals in the 1-2 ppm region, aliphatic signals at 2-3 ppm, and CH-N signals between 3 and 4 ppm; see Supplementary Fig. S6), were selected for further purification by MPLC (PuriFlash Column, 30µm, PF-30C18XS/55G; eluted with 70% to 100% MeOH in H₂O) to yield 1 (4.2 mg) and 2 (29.2 mg). Subfraction 4 W (288 mg) was fractionated by MPLC (PuriFlash Column, 30 µm, PF-30C18XS/55G; 60% to 100% MeOH in H_2O) to give six subfractions (4W1 to 4W6), of which 4W6 was found to possess signals characteristic of daphniphyllum alkaloids. Compound 4 (2.8 mg) was isolated from subfraction 4W6 by HPLC (65% MeOH in H₂O). Subfraction 4Z (2.0 g) was chromatographed on silica gel (1:1 to 0:1 hexane:EtOAc), followed by MPLC (PuriFlash Column, 30 µm, PF-30C18XS/55G; 70% to 100% MeOH in H₂O) to yield subfraction 4Z2A; the ¹H NMR signals of subfraction 4Z2A were found to be characteristic of daphniphyllum alkaloids. Further purification of subfraction 4Z2A by HPLC (65% MeOH in H₂O) yielded compound 3 (3.0 mg).

Glaulactam A (1): colorless oil; $[\alpha] + 102$ (*c* 0.42, MeOH); ECD (EtOH) λ_{max} ($\Delta \varepsilon$) 231 (-8.61), 270 (+12.81); UV (MeOH) λ_{max} (log ε) 260 (3.74), 270 (3.81), 280 (3.71) nm; IR (KBr) v_{max} 2949, 2933, 1736, 1730, 1712, 1695, 1433, 1371, 1247, 1224, 1195, 1168, 1028, and 754 cm⁻¹; ¹³C and ¹H NMR data, see Table 1; ESIMS *m/z* 486 [M + H]⁺, 508 [M + Na]⁺; HRESIMS *m/z* 486.2494 [M + H]⁺ (calcd. for C₂₇H₃₆NO₇, 486.2486).

Glaulactam B (2): colorless oil; $[\alpha]$ + 146 (*c* 0.58, MeOH); ECD (EtOH) λ_{max} ($\Delta \varepsilon$) 229 (-7.33), 270 (+5.40); UV (MeOH) λ_{max} (log ε) 261 (3.67), 271 (3.74), 280 (3.63) nm; IR (KBr) v_{max} 2949, 2931, 1734, 1699, 1685, 1436, 1375, 1363, 1244, 1193, 1166, 1028, and 752 cm⁻¹; ¹³C and ¹H NMR data, see Table 1; ESIMS *m*/*z* 428 [M + H]⁺, 450 [M + Na]⁺; HRESIMS *m*/*z* 428.2436 [M + H]⁺ (calcd. for C₂₅H₃₄NO₅, 428.2431).

Glaulactam C (3): colorless oil; $[\alpha] + 70$ (*c* 0.30, MeOH); ECD (EtOH) λ_{max} ($\Delta \varepsilon$) 226 (-9.02), 287 (+3.21); UV (MeOH) λ_{max} (log ε) 245 (4.07), 288 (3.91) nm; IR (KBr) ν_{max} 2926, 1735, 1701, 1687, 1641, 1597, 1436, 1365, 1284, 1234, 1195, 1172, 1118, 1062, 1028, and 754 cm⁻¹; ¹³C and ¹H NMR data, see Table 1; ESIMS *m/z* 448 [M + Na]⁺; HRESIMS *m/z* 448.2090 [M + Na]⁺ (calcd. for C₂₅H₃₁NO₅Na, 448.2094).

Cell culture and cytotoxicity assay. Madin-Darby canine kidney (MDCK) cells, (obtained from Dr. Lin, JH, Centers for Disease Control, Taipei, Taiwan) were cultured in Dulbecco's Modified Eagle's Medium (DMEM, Hyclone, GE Healthcare Life science) supplemented with 10% heat-inactivated fetal bovine serum (FBS, Hyclone, GE Healthcare Life science) and maintained at 37 °C in a 5% CO₂ atmosphere. For the cytotoxicity assay, MDCK cells were seeded in 96-well microplates (1×10^4 cells/well), and cultured for 24 h. Culture media were then replaced with media containing each of the indicated compounds for an additional 72 h. Cell viability was determined by the 3-(4,5-dimethylthiazol-2-yl)-5-(3- carboxymethoxyphenyl)-2-(4-sulfophenyl)-2H tetrazolium (MTS) assay kit purchased from Promega (Madison, WI).

Plaque-reduction assay. Monolayer MDCK cells were seeded in six-well plates (5×10^5 cells/well) for 24 h. The influenza virus A/WSN/33(H1N1) (100 plaque-forming units (PFU) per well) was mixed with each of the indicated compounds for 30 min at room temperature. The mixtures were subsequently adsorbed to the preseded cells for 1 h at 37 °C. After removal of the medium, the cells were washed with PBS (three times) and then overlaid with 0.3% agarose containing the indicated compounds for an additional 48 h at 37 °C, after which the cells were fixed with 10% formaldehyde for 1 h. Viral plaques were counted by staining with 0.5% crystal violet.

Mice and the preparation of bone marrow-derived murine dendritic cell (DC). According to the published method¹⁹, the murine bone-marrow-derived DCs were prepared from female C57BL/6 mice housed under controlled-temperature $(22 \pm 2 \,^{\circ}C)$ and humidity (45–65%) conditions, with a 12-h light/dark cycle and free access to food and water. The animals were treated according to the requirements of the Institutional Animal Care and Use Committee of National Chung-Hsing University.

Cell viability assay. DC-cell viability was measured using a cell-counting-kit assay (CCK-8; Dojindo Molecular Technologies, Inc., Kumamoto, Japan)²⁰, at an absorbance wavelength of 450 nm.

Cytokine and nitric oxide (NO) assay. After centrifugation at $1000 \times g$ for 15 min at 4 °C, the levels of tumor necrosis factor (TNF)- α (cat. no. 900-K54; PeproTech, Inc., London, UK), interleukin (IL)-6 (cat. no. 900-K50; PeproTech, Inc., London, UK), and IL-12p70 (cat. no. 900-K97; PeproTech, Inc., London, UK) in the culture supernatants were determined using murine ELISA kits for the respective cytokine according to the manufacturer's protocol. The levels of NO were measured indirectly by determining the concentration of NO²⁻ using a spectrophotometric assay based on the Griess reaction.

Computational details. Since different conformers of a specific stereochemical configuration can give different ECD spectra, it is critical to identify all relevant conformers in order to accurately predict the ECD spectrum. Therefore, standard conformational analyses were performed using the Confab program¹³. All optimized conformations were in a 10 kcal/mol energy window, with a root mean square (RMS) step-size of 0.2 Å. These conformers were then re-optimized at the B3LYP/6-31G(d) level of theory and verified to be true minima on the potential energy surface by frequency analyses. Then resulting geometries were subsequently used for three single-point calculations. CAM-B3LYP/TZVP²¹ calculations in the vacuum state were carried out in order to obtain converged wavefunctions for the ground states, which were used in the next two calculations. Energies were calculated at the M062x/Def2TZVP level with ethanol as the solvent, as this method provides more precise energies for conformational ordering²². The series of conformers was restricted by removing duplicates (RMSD < 0.2) and conformations outside of a 4 kcal/mol energy window, based on the Gibbs free energy at the M062x/Def2TZVP//B3LYP/6-31G(d) level in ethanol. The resulting structures were used in TDDFT ECD calculations, in ethanol as the solvent, at the TD-CAM-B3LYP/def2TZVP level by considering the 100 lowest-energy states. All calculation were performed using the Gaussian 09 program package with the "ultrafine grid" option, Integral (Grid = UltraFine), and all solvent effects were accounted for using the IEFPCM method^{14,23}. Finally, the overall ECD spectra were combined following Boltzmann weighting on the basis of the Gibbs free energies of the corresponding conformer calculated at the M062x/Def2TZVP//B3LYP/6-31G (d) level. All ECD spectra were processed with SpecDis and simulated by Gaussian functions with bandwidths (σ) of 0.16 eV and by considering velocity representations²⁴.

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Author Contributions

C.-H.C. and Y.-C.W. contributed equally to the experimental design and also analyzed the data. H.-C.H. analyzed the spectroscopic data. J.-C.C. and C.-C.L. performed the biological assays. T.P.G. and K.-W.H. calculated the ECD data. S.-Y.H. collected the samples and identified the species.

Additional Information

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