

Five new meroterpenoids from *Rhododendron anthopogonoides* and their anti-inflammatory activity

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Citation: Mengtian Li, Norbu Kelsang, Yongqin Zhao, Wensen Li, Feng Zhou, Pema , Lu Cui, Xianjie Bao, Qian Wang, Xin Feng, Minghua Yang, Five new meroterpenoids from *Rhododendron anthopogonoides* and their anti-inflammatory activity, *Chinese Journal of Natural Medicines*, 2025, 23(7), 881–887. doi: [10.1016/S1875-5364\(25\)60850-8](https://doi.org/10.1016/S1875-5364(25)60850-8).

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Original article

Five new meroterpenoids from *Rhododendron athopogonoides* and their anti-inflammatory activityMengtian Li^{a,Δ}, Norbu Kelsang^{b,c,Δ}, Yongqin Zhao^a, Wensen Li^a, Feng Zhou^{b,c}, Pema^{b,c}, Lu Cui^d, Xianjie Bao^d, Qian Wang^e, Xin Feng^{d,e,*}, Minghua Yang^{a,*}^a State Key Laboratory of Natural Medicines, Jiangsu Key Laboratory of Bioactive Natural Product Research, School of Traditional Chinese Pharmacy, China Pharmaceutical University, Nanjing 210009, China^b Tibet Ganlu Pharmaceutical Technology Company, Lhasa 850032, China^c Technological Innovation Center of Traditional Tibetan Medicine Modernization of Tibet Autonomous Region, Lhasa 850032, China^d Beijing Hospital of Tibetan Medicine, China Tibetology Research Center, Beijing 100029, China^e Tibetan Medicine Institute, China Tibetology Research Center China, Beijing 100101, China

ARTICLE INFO

Article history:

Received 23 August 2024

Revised 16 October 2024

Accepted 23 December 2024

Available online 20 July 2025

Keywords:

Anti-inflammatory

Tibetan medicine

Natural products

Isolation and identification

Rhododendron athopogonosides Maxim

ABSTRACT

Five meroterpenoids, rhodonoids K–M (1–2), daurichromene E (3), and grifolins A–B (4–5), together with seven known compounds (6–12), were isolated from *Rhododendron athopogonoides*. The chemical structures of these compounds were elucidated through comprehensive analysis of high-resolution electrospray ionization mass spectrometry (HR-ESI-MS), ultraviolet (UV), infrared spectroscopy (IR), and nuclear magnetic resonance (NMR) data. Their absolute configurations were determined by comparing experimental electronic circular dichroism (ECD) spectra with computed values. Notably, compounds 1 and 3 demonstrated significant inhibitory effects on lipopolysaccharide (LPS)-induced inflammation in RAW264.7 cells. These compounds markedly suppressed the mRNA expressions of inflammatory factors, including interleukin (IL)-1 β , IL-6, and tumor necrosis factor- α (TNF- α) while also down-regulating the protein expressions of inducible nitric oxide synthase (iNOS) and cyclooxygenase-2 (COX-2).

1. Introduction

Rhododendron athopogonosides Maxim., a perennial evergreen shrub, thrives in high-altitude regions across several provinces in China, including Sichuan, Gansu, Yunnan Provinces, and the Tibet Autonomous Region^{1,2}. It is a well-documented medicinal species in Tibetan ethnomedicine, where its flowers, leaves, and twigs are traditionally employed to treat disorders collectively referred to as “Peigen” in the Tibetan language³, making *R. athopogonosides* a prevalent component in traditional Tibetan prescriptions. Phytochemical investigations have revealed that the extracts of *R. athopogonosides* exhibit various bioactivities, such as antioxidant⁴, insecticidal², bacteriostatic⁵, anti-inflammatory activities⁶. These biological properties have driven increasing interest in identifying its pharmacologically active constituents, which are primarily classified as flavonoids⁷, essential oils⁸, and triterpenes⁹. Among these, flavonoids and essential oils have been extensively studied and are recognized for their therapeutic relevance in respiratory and cardiovascular diseases¹⁰.

Meroterpenoids, a structurally diverse class of natural

products resulting from hybrid biosynthetic pathways, are typically categorized into polyketide-terpenoid and non-polyketide terpenoid subtypes¹¹ based on their biosynthetic origin. The genus *Rhododendron* has emerged as a prolific source of polyketide-derived meroterpenoids¹², especially those containing chromane or chromene scaffolds. This genus has yielded over one hundred meroterpenoids^{13,14} exhibiting diverse structural skeletons and biological activities. Although *R. athopogonosides* is known to produce meroterpenoids, systematic studies of this species remain relatively scarce. Iwata and Kitanaka¹⁵, followed by Shi et al.⁶, reported several structurally novel meroterpenoids from this plant, highlighting their significant anti-inflammatory potential. Driven by our continuing interest in bioactive meroterpenoids from medicinal plants^{16,17}, we investigated the chemical composition of *R. athopogonosides* from Tibet. This work led to the isolation and structural characterization of five previously undescribed meroterpenoids (1–5) (Fig. 1), and seven known compounds (6–12). Among these, compounds 1 and 3 exhibited significant anti-inflammatory effects in lipopolysaccharide (LPS)-induced RAW264.7 cells. This study presents the isolation, structural elucidation, and anti-inflammatory activity of these compounds.

2. Results and discussion

Compound 1 was obtained as a yellow oil. Its molecular for-

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mula $C_{22}H_{30}O_3$ was established by the high-resolution electrospray ionization mass spectrometry (HR-ESI-MS) (m/z 365.2089 $[M + Na]^+$, Calcd. for $C_{22}H_{30}O_3Na$ 365.2087), indicating eight degrees of structural unsaturation. The 1H nuclear magnetic resonance (NMR) spectrum (Table 1) showed two phenyl protons [δ_H 6.16 (s, H-6); 6.28 (s, H-8)], one olefinic proton [δ_H 5.46 (t, $J = 5.4$ Hz, H-15)], a pair of terminal olefinic protons [δ_H 4.80 (s, H-19a); 4.58 (s, H-19b)], and three methyls [δ_H 1.70 (s, H-18); 1.35 (s, H-20); 2.20 (s, H-21)]. Heteronuclear single quantum coherence (HSQC) correlations facilitated the identification of additional protons, including seven methylenes [δ_H 1.89 (m, $J = 13.2$, 3.0 Hz, H-3); 1.54 (m, H-9); 1.57 (m, H-10); 2.27 (m, H-13); 2.28 (m, H-14); 4.01 (s, H-17)] and three methines [δ_H 3.25 (m, H-4); 2.29 (m, H-11); 5.46 (t, H-15)]. The corresponding carbons were all found in the ^{13}C NMR spectrum (Table 1), which also included six non-proton-bearing olefinic ones and one quaternary carbon. These data were similar to those of rhodonoid G¹⁸, a known meroterpenoid from *R. athopogonosides*.

The identical chromane core was confirmed by the similar chemical shifts of downfield-shifted C-2 and C-8a, along with the heteronuclear multiple bond correlations (HMBCs) (Fig. 2), especially from H-4 to C-2/C-3/C-8a, from H-6 to C-4a/C-5, from H-8 to C-4a/C-8a, from H₃-20 to C-2/C-3, and from H₃-21 to C-6/C-7/C-8. The correlations from H-9 to C-2/C-3/C-10/C-11, from H-4 to C-4a/C-5, and from H-20 to C-9 further established the same benzo[*c*]-2-oxabicyclo[3.3.1]nonane structure like rhodonoid G, which was also consistent with the observed spin system from C-3 to C-9 in 1H - 1H correlation spectroscopy (COSY) spectrum (Fig. 2). The difference was found at the substitution of C-11 in **1**. Compound **1** exhibited a 2-methylhepta-2,6-dien-1-ol side chain, evidenced by the 1H - 1H COSY correlations (Fig. 2) of H-13/H-14/H-15 and the HMBCs (Fig. 2) from H-18 to C-15/C-16/C-17, from H-15 to C-13/C-14/C-17, and from H-19 to C-11/C-12/C-13. It was settled at C-11 by the HMBCs (Fig. 2) from H-13 to C-11 and C-14. Thus, its planner structure was determined as shown. In the ROESY experiment (Fig. 3), the correlation between H-15 and H-17 indicated the *E*-orientation of C-15/C-16 double bond. The correlations from H-3a to H-4 and H₃-20 revealed their identical orientation. H-11, however, showed no useful correlations for configuration determination. The theoretical calculation of electronic circular dichroism (ECD) spectra was conducted to determine the configuration, which was based on time-dependent density functional theory (TD-DFT) at the B3LYP/6-31 + G(d, p) level. The two possible combinations of C-11 configuration were both submitted for calculation. The calculated ECD curves of

2*S*,4*S*,11*R* showed an excellent fit with the experimental one. Thus, the structure of compound **1** was determined as shown.

Compound **2**, a brown oil, was assigned a molecular formula of $C_{16}H_{20}O_3$ by HR-ESI-MS (m/z 261.1485 $[M + H]^+$, Calcd. for $C_{16}H_{21}O_3$ 261.1485), indicating seven degrees of unsaturation. The 1D NMR data of **2** resembled to those of **1** (Table 2), and the same benzo[*c*]-2-oxabicyclo[3.3.1]nonane ring system was further determined by the similar 1H - 1H COSY and HMBCs (Fig. 2). The key structural difference was the presence of an acetyl group at C-11, as evidenced by HMBCs (Fig. 2) from methyl H-13 to C-11 and carbonyl C-12 and from H-10 and H-4 to C-12. In addition, rotating frame nuclear Overhauser effect spectroscopy (ROESY) revealed a consistent spatial orientation of H-4 and H₃-14, while the lack of correlation between H-4 and H-11 indicated an opposite orientation at C-11. ECD calculations (Fig. 4) confirmed the absolute configuration as 2*S*,4*S*,11*R* configuration for **2**.

Compound **3** was isolated as a brown oil with molecular formula $C_{22}H_{32}O_4$ determined by the HR-ESI-MS (m/z 383.2192 $[M + Na]^+$, Calcd. for $C_{22}H_{32}O_4Na$ 383.2193). The NMR data of **3** showed high similarity with daurichromene D¹⁹, a prenylated orcinol derivative isolated from *R. dauricum*. However, compound **3** exhibited one additional oxygen and two additional hydrogen atoms, indicating that **3** is an additional product of daurichromene D¹⁹. This inference was corroborated by the absence of C-3/C-4 double bond signals and the presence of an oxygen-bearing methenyl at δ_C 68.3. Further evidence was provided by the 1H - 1H COSY correlations (Fig. 2) between H-3 and H-4, and the HMBCs (Fig. 2) from H-4 to C-2, C-3, C-4a, C-8a and C-5, and from H-3 to C-20. The comprehensive analysis of 2D NMR data further determined the planar structure of **3**. In ROESY spectrum (Fig. 3), the correlations of H-11/H-13 and H-15/H-17 established the *E*-orientation of double bonds in the side chain. The correlation of H-3/H-20 indicated their same orientation. To elucidate the absolute configuration, time-dependent density functional theory calculations of ECD spectra were performed (Fig. 4). The experimental spectrum closely aligned with the calculated one for the 2*R*,3*R* configurations. Thus, **3** was determined as shown.

Compound **4**, obtained as a brown oil, had a molecular formula of $C_{22}H_{32}O_3$ based on HR-ESI-MS ion at m/z 367.2244 $[M + Na]^+$ (Calcd. for $C_{22}H_{32}O_3Na$ 367.2244). Its 1H NMR spectrum (Table 3) showed signals for two *meta*-coupled aromatic protons [δ_H 6.22 (s, H-4); 6.22 (s, H-6)], three olefinic protons [δ_H 5.25 (t, $J = 6.5$ Hz, H-8); 5.07 (t, $J = 6.5$ Hz, H-12); 5.38 (t, $J = 6.5$ Hz, H-16)], and four methyls [δ_H 2.18 (s, H-22); 1.66 (s, H-19); 1.58 (s, H-20); 1.80 (s, H-21)]. The corresponding carbons were also found in ^{13}C

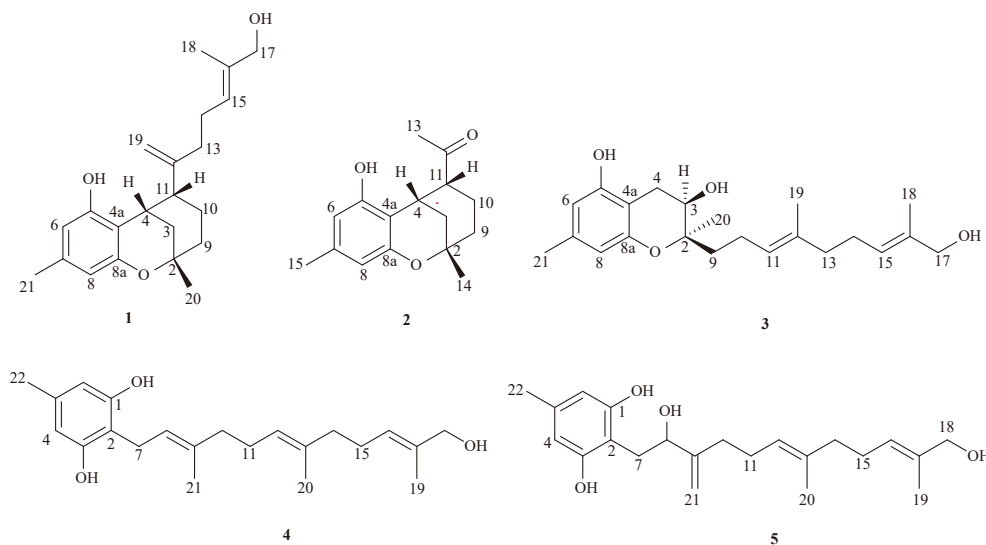


Fig. 1 Structures of compounds 1-5.

Table 1 The ^1H (600 MHz) and ^{13}C (150 MHz) NMR data for **1** and **2** (J in Hz).

No.	1 ^a		2 ^b	
	δ_{H} (J in Hz)	δ_{C}	δ_{H} (J in Hz)	δ_{C}
2		73.8		74.6
3	1.89, dd (13.2, 3.0)	37.6	2.00, dd (13.2, 3.0)	38.0
	1.79, dt (13.2, 3.0)		1.76, dt (12.6, 3.0)	
4	3.25, m	32.2	3.74, br s	30.7
4a		107.8		107.4
5		153.4		155.9
6	6.16, s	108.3	6.04, s	107.4
7		137.5		138.8
8	6.28, s	109.0	6.09, s	108.2
8a		157.2		158.6
9	1.54, m	39.8	1.93, dt (9.0, 2.4)	39.6
	2.01, m		1.56, m	
10	1.57, m	23.5	1.57, m	20.9
	1.42, m		1.41, m	
11	2.29, m	47.6	2.83, m	56.1
12		152.0		213.6
13	2.27, m	35.7	2.28, s	29.1
	2.17, m			
14	2.28, m	25.0	1.33, s	29.2
15	5.46, t (5.4)	125.4	2.15, s	21.4
16		135.8		
17	4.01, s	68.8		
18	1.70, s	13.9		
19	4.58, s	110.0		
	4.80, s			
20	1.35, s	28.8		
21	2.20, s	21.4		

^a in CDCl_3 , ^b in CD_3OD .

NMR. Additionally, HSQC correlations indicated the presence of six methylenes. These 1D NMR data were comparable to those of grifolin²⁰, while the presence of a downfield-shifted methylene (δ_{C} 69.1) revealed an oxidative methyl in **4**. The singlet methylene at δ_{H} 4.00 showed HMBs with C-16 and C-19 (Fig. 2), confirming the hydroxyl substitution at C-18. Furthermore, ROESY correlations from H-8 to H-10, from H-12 to H-14, and from H-16 to H-18 indicated the *trans*-configuration of the three double bonds (Fig. 3).

Compound **5**, also a brown oil, had a molecular formula of $\text{C}_{22}\text{H}_{32}\text{O}_4$ (m/z 383.2193 [$\text{M} + \text{Na}$]⁺, Calcd. 383.2193), indicating an additional oxygen atom compared to **4**. Structural elucidation using 2D NMR revealed that **5** is an oxidized derivative of **4**. Key HMBs from terminal olefinic protons [δ_{H} 4.88 (m, H-21a); 5.10 (d, $J = 7.0$ Hz, H-21b)] (Fig. 2) to C-8, C-9, and C-10 were detected, and the methine proton at C-8 [δ_{H} 4.31 (d, $J = 6.5$ Hz, H-8)] displayed ^1H - ^1H COSY correlation (Fig. 2) with H-7 and critical HMBs (Fig. 2) with C-2, C-7, C-10, and C-21. Additional 2D NMR data (Fig. 2) further confirmed the structure of **5**. The near-zero optical rotation and flat ECD curve suggested a racemic or nearly

racemic mixture, which was validated by chiral high-performance liquid chromatography (HPLC), revealing two enantiomeric peaks in approximately equal proportions.

The remaining known compounds were identified as loliolide (**6**)²¹, caffeic acid methyl ester (**7**)²², chlospicate C (**8**)²³, istanbulin A (**9**)²⁴, 4 α -hydroxy-5 α ,8 β (*H*)-eduesm-7(11)-en-8,12-olidel (**10**)²⁵, chlospicate D (**11**)²³, and 6(1*H*)-azulenone (**12**)²⁶ by comparing their NMR data with those reported in literature.

All compounds were evaluated for cytotoxicity and anti-inflammatory activity in LPS-induced RAW264.7 macrophages. The CCK-8 assay confirmed that all tested compounds were non-toxic at concentrations up to 50 $\mu\text{mol}\cdot\text{L}^{-1}$. The compounds were then tested for their inhibitory effects on NO production, with L-NMMA (IC_{50} 40.58 \pm 0.86 $\mu\text{mol}\cdot\text{L}^{-1}$) as the positive control. Most compounds showed weak or no inhibition (IC_{50} 50 $\mu\text{mol}\cdot\text{L}^{-1}$), whereas compounds **1** and **3** displayed moderate NO inhibition with IC_{50} values of 10.75 \pm 2.28 and 15.85 \pm 1.58 $\mu\text{mol}\cdot\text{L}^{-1}$, respectively. To explore their mechanism of action, the effects of compounds **1** and **3** on the mRNA expression of key inflammatory markers—tumor necrosis factor- α (*TNF- α*), interleukin (*IL*)-1 β , *IL*-6, cyclooxygenase-2 (*COX*-2), and inducible nitric oxide synthase (*iNOS*)—were assessed by quantitative reverse transcription-polymerase chain reaction (qRT-PCR). As shown in Figs. 5A–5E, Both compounds significantly suppressed these markers in a dose-dependent manner. They both showed a good inhibition rate of *IL*-1 β since about 50% mRNA expression was inhibited at the concentration of 12.5 $\mu\text{mol}\cdot\text{L}^{-1}$. Compound **1** exhibited stronger suppression of *COX*-2 mRNA (34.5% \pm 3.22% inhibition), while compound **3** more effectively reduced *iNOS* expression. Western blotting analysis confirmed these effects at the protein level (Figs. 5F–5K): compound **1** preferentially downregulated *COX*-2, while **3** was more effectively suppressed *iNOS* expression, suggesting divergent but complementary anti-inflammatory mechanisms.

3. Conclusion

In this study, five previously undescribed meroterpenoids, rhodonoids K and M (**1** and **2**), daurichromene E (**3**), and grifolins A and B (**4** and **5**), were isolated and structurally elucidated from *R. anthopogonoides*. Their structures and absolute configurations were determined through comprehensive spectroscopic analysis. Compounds **1** and **3** exhibited moderate but significant anti-inflammatory activity in LPS-stimulated RAW264.7 macrophages, demonstrated by their ability to reduce nitric oxide production and downregulate the expression of key inflammatory mediators at both mRNA and protein levels. Thus, these meroterpenoids are considered to be the bioactive constituents responsible for the anti-inflammatory properties of *R. anthopogonoides*.

4. Experimental

4.1. Materials and methods

ECD spectra were obtained using a Jasco J-810 Circular dichroism spectrometer (JASCO, Japan). Ultraviolet (UV) spectra were obtained using a UV-2450 spectrophotometer (Shimadzu, Japan). IR spectra were measured using a Bruker Tensor-27 spectrophotometer (Bruker, Germany). Optical rotations were measured on a Jasco P-1020 polarimeter (JASCO, Japan). NMR spectra were recorded on a Bruker Avance III-500 or Bruker Avance III-600 spectrometer (Bruker, Germany) with TMS as an internal standard. HR-ESI-MS data were recorded on Micro Q-TOF MS spectrometer (Waters, USA). ESI-MS data were recorded on Agilent 1100 (Agilent, USA). Column chromatography was performed on silica gel (200–300 mesh, 100–200 mesh, Qingdao

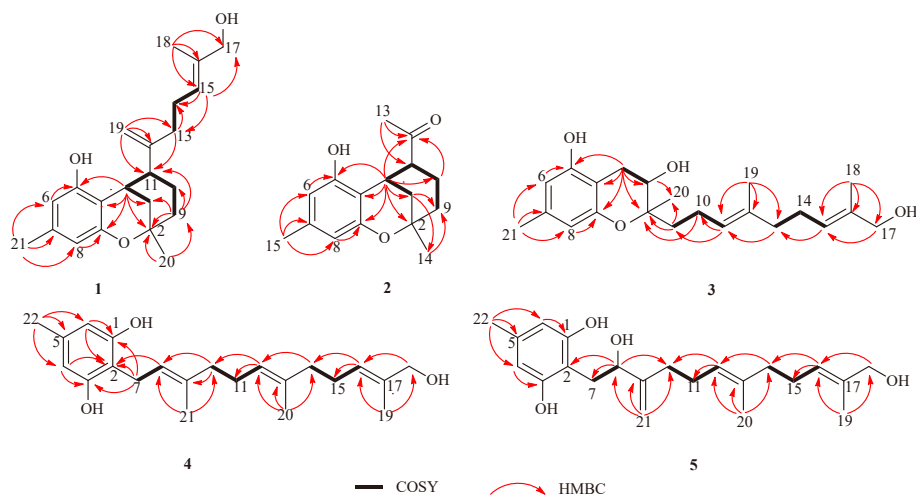


Fig. 2 Key ^1H - ^1H COSY and HMBCs of compounds 1-5.

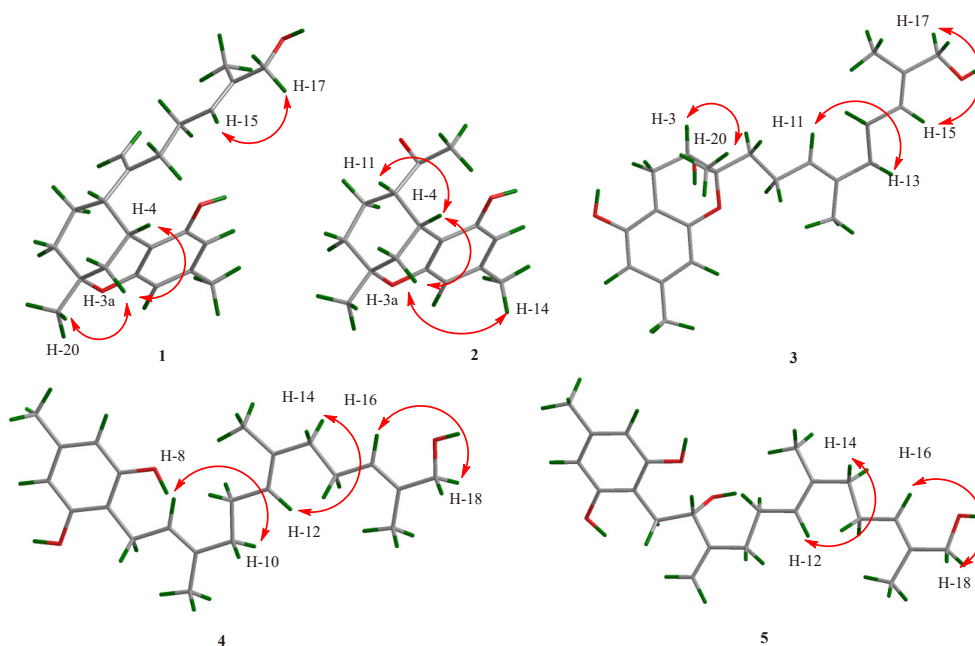


Fig. 3 Key ROSEY correlations of compounds 1-5.

Table 2 The ^1H (600 MHz) and ^{13}C (125 MHz) NMR data for 3 in CDCl_3 (J in Hz).

No.	δ_{H}	δ_{C}	No.	δ_{H}	δ_{C}
2		78.4	11	5.12, t (6.0)	122.4
3	3.87, t (4.5)	68.3	12		132.7
4	2.62, dd (14.0, 5.0)	28.3	13	2.02, t (6.0)	41.1
	2.87, dd (14.0, 5.0)		14	2.12, m	28.3
4a		102.6	15	5.34, t (5.0)	123.5
5		150.0	16		132.3
6	6.19, s	106.5	17	3.98, s	69.4
7		135.1	18	1.64, s	16.8
8	6.27, s	108.6	19	1.59, s	18.9
8a		150.8	20	1.28, s	22.0
9	1.64, m	39.0	21	2.20, s	24.0
10	2.12, m	24.4			

Marine Chemical Inc., Qingdao, China), MCI gel CHP20/P120 (37–73 μm , Mitsubishi Chemical Industries, Tokyo, Japan), RP-18 (40–63 μm , Fuji Silysia Chemical Ltd., Tokyo, Japan), and Sephadex LH-20 (Amersham Pharmacia, Uppsala, Sweden). Preparative HPLC was carried out by using a Shimadzu LC-8A chromatograph equipped with a Shim-pack ODS (250 mm \times 20 mm, 5 μm). Analytical HPLC was performed using an Agilent 1260 chromatograph equipped with a Shim-pack ODS (4.6 mm \times 250 mm, 5 μm).

4.2. Plant material

R. anthopogonoides specimens were collected from Tibet, China in August 2021. The plant material was taxonomically identified by Prof. Chaofeng Zhang of China Pharmaceutical University. The voucher specimen (No. 2022-LXDJ) was deposited at the Herbarium of the Department of Chemistry, China Pharmaceutical University (China).

4.3. Extraction and isolation

Dried and powdered aerial parts of *R. anthopogonoides* (1 kg)

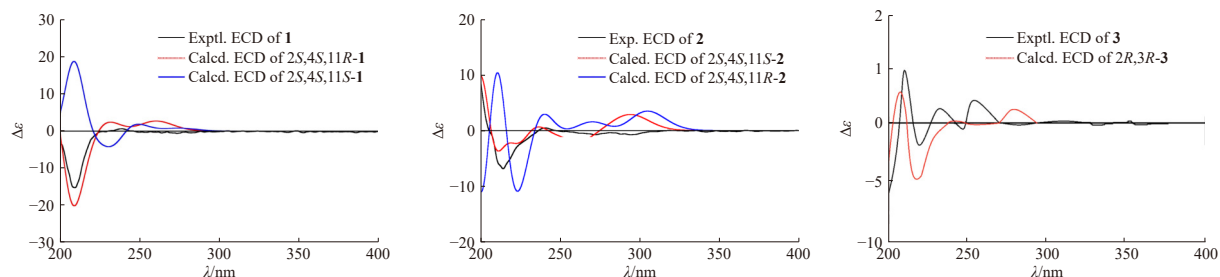


Fig. 4 Experimental and calculated ECD of compounds 1-3.

Table 3 The ^1H (600 MHz) and ^{13}C (125 MHz) NMR data for **4** and **5** in CDCl_3 (J in Hz)

No.	4		5	
	δ_{H} (J in Hz)	δ_{C}	δ_{H} (J in Hz)	δ_{C}
1		155.0		155.7
2		110.8		110.4
3		155.0		155.7
4	6.22, s	109.0	6.26, s	109.4
5		137.4		138.0
6	6.22, s	109.0	6.26, s	109.4
7	3.38, d (6.5)	22.3	2.73, dd (12.0, 7.5) 3.09, d (12.0)	29.8
8	5.25, t (6.5)	122.2	4.31, d (6.5)	77.0
9		138.2		151.4
10	2.03, m	39.7	2.20, m	26.1
11	2.06, m	26.2	2.19, m	32.2
12	5.07, t (6.5)	126.7	5.13, m	124.7
13		135.1		134.8
14	2.07, m	39.2	2.05, m	39.0
15	2.08, m	26.0	2.14, m	25.5
16	5.38, t (6.5)	126.7	5.33, m	125.7
17		134.3		134.3
18	4.00, s	69.1	3.98, s	68.8
19	1.66, s	13.8	1.63, s	13.9
20	1.58, s	16.1	1.60, s	16.1
21	1.80, s	16.3	4.88, m 5.10, d (7.0)	108.9
22	2.18, s	21.2	2.20, s	21.3

were refluxed with 95% EtOH (3×5 L, 3 h each). The combined extracts were concentrated under reduced pressure to yield approximately 100 g of residue, which was then suspended in water and partitioned thrice with petroleum ether (PE) and ethyl acetate (EtOAc) sequentially. The PE fraction (20 g) was subjected to silica gel column chromatography (elution with PE/EtOAc, gradient from 1:0 to 1:2, V/V), yielding five fractions (Frs. 1-5). Fr. 3 (2.5 g) was further purified using a Sephadex LH-20 column eluted with methanol (MeOH)/dichloromethane (CH_2Cl_2) (1:1, V/V), yielding 8 subfractions (Frs. 3.1-3.8). Fr. 3.2 (210.8 mg) was then subjected to preparative HPLC (MeOH- H_2O , 40:60, V/V) to yield compounds **1** (6.4 mg, t_{R} 37.5 min) and **2** (9.2 mg, t_{R} 42.6 min). Fr. 5 (3.5 g) was subjected to an ODS column eluted

with MeOH/ H_2O (35:65-100:0, V/V) to obtain 12 subfractions (Frs. 5.1-5.12). Fr. 5.2 (202.4 mg) was further purified by preparative HPLC (MeOH- H_2O , 60:40, V/V) to yield compound **3** (10.1 mg, t_{R} 29.4 min). Fr. 5.4 (222.8 mg) was also purified by preparative HPLC (MeOH- H_2O , 50:50, V/V) to obtain compounds **4** (11.0 mg, t_{R} 38.4 min) and **5** (8.6 mg, t_{R} 40.2 min). The EtOH part (40 g) was separated using a silica gel column and eluted with $\text{CH}_2\text{Cl}_2/\text{MeOH}$ (1:0-1:2, V/V) to yield 8 fractions (Frs. 6-13). Fr. 8 (4.0 g) was further purified using an ODS column eluted with MeOH/ H_2O (35:65-100:0, V/V), yielding 14 subfractions (Frs. 8.1-8.14). Fr. 8.2 (120.5 mg) was finally purified by preparative HPLC (MeOH- H_2O , 45:55, V/V) to obtain compound **6** (20.4 mg, t_{R} 20.4 min). Compounds **7** (15.6 mg, t_{R} 21.0 min) and **8** (9.5 mg, t_{R} 24.3 min) were isolated from Fr. 8.5 using preparative HPLC (MeOH- H_2O , 45:55, V/V). Fr. 8.4 was finally purified by preparative HPLC (MeOH- H_2O , 45:55, V/V) to obtain compound **9** (8.7 mg, t_{R} 30.6 min). Fr. 8.5 placed steadily to precipitate compound **10** (16.8 mg). And finally, Fr. 8.6 was purified by preparative HPLC (MeOH- H_2O , 50:50, V/V) to obtain compounds **11** (11.3 mg, t_{R} 37.5 min) and **12** (9.3 mg, t_{R} 38.2 min).

Rhodonoid K: yellow oil; $[\alpha]_{\text{D}}^{25} +16.0$ (c 0.1, MeOH); UV (MeOH) λ_{max} ($\log \epsilon$) 282 (3.03), 274 (3.04), 208 (4.15), 193 (3.58) nm; IR (KBr) ν_{max} 3333, 2930, 1625, 1586, 1451, 1383, 1160, 1134, 1094, 997, 883 cm^{-1} ; ^1H and ^{13}C NMR data (Table 1); HR-ESI-MS m/z 365.2089 $[\text{M} + \text{Na}]^+$ (Calcd. for $\text{C}_{22}\text{H}_{30}\text{O}_3\text{Na}$, 365.2087).

Rhodonoid L: brown oil; $[\alpha]_{\text{D}}^{25} -20.8$ (c 0.1, MeOH); UV (MeOH) λ_{max} ($\log \epsilon$) 274 (3.09), 207 (3.96), 193 (3.48) nm; IR (KBr) ν_{max} 3269, 2931, 1686, 1624, 1588, 1451, 1347, 1159, 1097, 997, 823 cm^{-1} ; ^1H and ^{13}C NMR data (Table 1); HR-ESI-MS m/z 261.1485 $[\text{M} + \text{H}]^+$ (Calcd. for $\text{C}_{16}\text{H}_{21}\text{O}_3$, 261.1485).

Daurichromene E: brown oil; $[\alpha]_{\text{D}}^{25} -9.6$ (c 0.1, MeOH); UV (MeOH) λ_{max} ($\log \epsilon$) 280 (2.87), 273 (2.89), 207 (4.16) nm; IR (KBr) ν_{max} 3374, 2922, 1628, 1589, 1518, 1452, 1383, 1208, 1073, 996, 823, 574 cm^{-1} ; ^1H and ^{13}C NMR data (Table 2); HR-ESI-MS m/z 383.2192 $[\text{M} + \text{Na}]^+$ (Calcd. for $\text{C}_{22}\text{H}_{32}\text{O}_4\text{Na}$, 383.2193).

Grifolin A: brown oil; UV (MeOH) λ_{max} ($\log \epsilon$) 276 (2.73), 207 (4.13) nm; IR (KBr) ν_{max} 3404, 2922, 1715, 1629, 1597, 1451, 1385, 1222, 1049, 989, 824, 591 cm^{-1} ; ^1H and ^{13}C NMR data (Table 3); HR-ESI-MS m/z 367.2242 $[\text{M} + \text{Na}]^+$ (Calcd. for $\text{C}_{22}\text{H}_{32}\text{O}_3\text{Na}$, 367.2244).

Grifolin B: brown oil; $[\alpha]_{\text{D}}^{25} -1.4$ (c 0.1, MeOH); UV (MeOH) λ_{max} ($\log \epsilon$) 276 (2.88), 207 (4.23) nm; IR (KBr) ν_{max} 3343, 2923, 1716, 1633, 1591, 1451, 1386, 1210, 1073, 903, 826, 590 cm^{-1} ; ^1H and ^{13}C NMR data (Table 3); HR-ESI-MS m/z 367.2242 $[\text{M} + \text{Na}]^+$ (Calcd. for $\text{C}_{22}\text{H}_{32}\text{O}_3\text{Na}$, 367.2244).

4.4. Cell culture and viability assay

RAW264.7 cells were cultured in high-glucose DMEM supplemented with 10% FBS and 1% penicillin-streptomycin antibiotics in the atmosphere with 5% CO_2 at 37 $^\circ\text{C}$. The isolated compounds were dissolved in DMSO for activity testing. Cell viability was assessed by cell counting kit-8 (CCK-8) assay. The cells (5×10^3 cells/well) were cultured in 96-well plates for 12 h, fol-

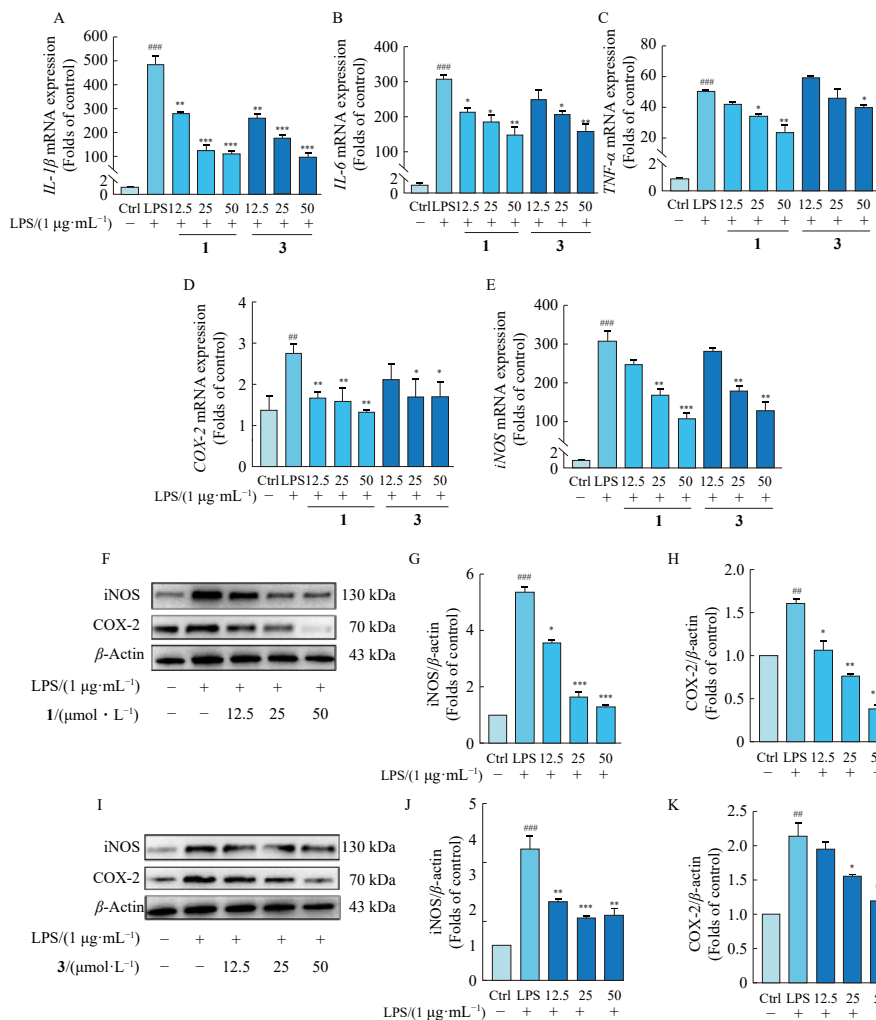


Fig. 5 Effects of compounds **1** and **3** on LPS-induced mRNA expression of *IL-1β* (A), *IL-6* (B), *TNF-α* (C), *COX-2* (D), and *iNOS* (E) in RAW264.7 cells were determined by qRT-PCR. Effects of compounds **1** (F–H) and **3** (I–K) on LPS-induced protein expression of COX-2 and iNOS in RAW264.7 cells were determined by Western blotting. Data are presented as mean \pm SD ($n = 3$). * $P < 0.05$, ** $P < 0.01$ and *** $P < 0.001$ vs control group, # $P < 0.05$, ## $P < 0.01$ and ### $P < 0.001$ vs LPS group.

lowed by treatment with the tested compounds **1–12** separately at different concentrations (0, 6.25, 12.5, 25, 50, 100 $\mu\text{mol}\cdot\text{L}^{-1}$) for 24 h. After incubation with CCK-8 reagent (20 μL) for 1 h, the absorbance at 450 nm was measured with a microplate reader (TECAN, Switzerland).

4.5. NO production assay

NO levels in the culture supernatant were quantified using the Griess reagent nitrite measurement kit (Beyotime, China). RAW264.7 cells (2×10^5 cells/well) were cultured in 100 μL DMEM in 96-well plates for 12 h and then treated with different concentrations (0, 6.25, 12.5, 25, 50 $\mu\text{mol}\cdot\text{L}^{-1}$) of compounds **1–5** for an additional 12 h. Lipopolysaccharide (LPS) at 1 $\mu\text{g}\cdot\text{mL}^{-1}$ was added as an inducer and cultured together with the compounds. After incubation, 50 μL of the supernatant was mixed with an equal volume of Griess reagent A and B. Following a 10-min shaking period, the absorbance was measured at 450 nm using a microplate reader (TECAN, Switzerland), and subsequently, the NO inhibition rates were calculated.

4.6. Quantitative real-time PCR analysis

Total RNA was isolated using a total RNA isolation kit (Vazyme, China), and RNA concentration was measured using a NanoDrop One Spectrophotometer (Thermo, USA). First-strand cDNA was synthesized from total RNA and used for qRT-PCR us-

ing SYBR Green Master Mix (Vazyme, China) on a Roche Light-Cycler system. Expression levels of *TNF-α*, *IL-1β*, *IL-6*, *COX-2*, and *iNOS* were quantitatively analyzed using the Real-Time PCR Detection System (Roche, Switzerland). The quantitative primers employed for amplification are listed in Supplementary Table 1.

4.7. Western blotting analysis

RAW264.7 cells (1×10^5 cells/well) were cultured in 6-well plates for 12 h and pretreated with compounds **1** or **3** at gradient concentrations for 12 h, followed by co-stimulation with 1 $\mu\text{g}\cdot\text{mL}^{-1}$ LPS. Total protein was extracted and quantified using a bicinchoninic acid (BCA) assay. Equal amounts of protein were separated by sodium dodecyl sulfate-polyacrylamide gel electrophoresis (SDS-PAGE) and transferred onto the immunoblot membranes. Protein bands were detected using an enzyme marker. Membranes were blocked and incubated overnight at 4 $^{\circ}\text{C}$ with primary antibodies against COX-2, iNOS, or β -actin. After washing with tris-buffered saline with Tween 20 (TBST), membranes were incubated with secondary antibodies at room temperature for 1 h. Finally, the membranes were visualized using a Gel Documentation System.

Funding

This work was supported by the second batch of Bureau-level Projects of the Tibetan Medicine Administration of the Tibet

Autonomous Region in 2020 (No. JJKT20200024).

Availability of supporting information

Supporting information for this study can be obtained by contacting the corresponding authors *via* E-mail.

Declaration of competing interest

These authors have no conflict of interest to declare.

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