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3,4-Seco-Dammarane Triterpenoid Saponins with Anti-inflammatory Activity Isolated from the Leaves of Cyclocarya paliurus

Wei Liu, Shengping Deng, Dexiong Zhou, Yan Huang, Chenguo Li, Lili Hao, Gaorong Zhang, Shanshan Su, Xia Xu, Rui-Yun Yang, Jun Li, and Xishan Huang

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1	3,4-Seco-Dammarane Triterpenoid Saponins with Anti-inflammatory
2	Activity Isolated from the Leaves of Cyclocarya paliurus

3 Wei Liu, Shengping Deng, Dexiong Zhou, Yan Huang, Chenguo Li, Lili Hao, Gaorong Zhang, 4 Shanshan Su, Xia Xu, Ruiyun Yang, Jun Li*, Xishan Huang*

5 State Key Laboratory for the Chemistry and Molecular Engineering of Medicinal Resources,

6 School of Chemistry and Pharmaceutical Sciences of Guangxi Normal University, Guilin 541004, 7

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China

9 ABSTRACT: Cyclocarya paliurusis commonly used for the prevention and treatment 10 of hypertension, diabetes, and inflammation in South China. Although research on the anti-inflammatory effects of C. paliurus leaves has been reported, no active 11 12 anti-inflammatory compounds have been identified. In the present study, RAW 264.7 cells were used to establish a bioactivity-guided identification model to verify the 13 inhibitory effects of C. paliurus leaves on inflammation and identify the 14 anti-inflammatory constituents. The active fraction was isolated to yield eighteen 15 16 dammarane triterpenoid saponins, including eleven new 3,4-seco-dammarane 17 triterpenoid saponins (1-11), the structures of which were identified on the basis of 18 analyses of nuclear magnetic resonance (NMR) spectroscopy, mass spectrometry (MS) and comparison with literature data. Compounds 7, 8, 10, and 11 showed strong 19 20 inhibition on nitric oxide (NO) productions, with IC₅₀ values ranging from 8.23 μ M to 21 11.23 μ M. These four compounds significantly decreased the secretion of tumor 22 necrosis factor-alpha (TNF- α), prostaglandin E₂ (PGE₂) and interleukin 6 (IL-6) in 23 lipopolysaccharide (LPS)-activated RAW 264.7 cells. Furthermore, compound 7 24 decreased the expression of the proteins cyclooxygenase-2 (COX-2), inducible nitric-oxide synthase (iNOS), and nuclear factor kappa-B (NF- κ B/p65). In addition, 25

the structure-activity relationships of the isolates were investigated. The results suggest that 3,4-*seco*-dammarane triterpenoid saponins may be used as potential anti-inflammatory drugs and warrant further investigation.

29

30 INTRODUCTION

Cyclocarya paliurus (Batal.) Iljinskaja, a Chinese indigenous plant, is the sole 31 32 species in the genus Cyclocarya Iljinskaja of the Juglandaceae family. The leaves of C. paliurus are used not only as drug formulations in traditional Chinese medicine to 33 34 treat hyperlipidemia, hypertension, diabetes mellitus, chronic inflammatory diseases and as an antioxidant but also as a famous "sweet tea", with a very sweet taste, in folk 35 medicine in China to prevent and treat hypertension, diabetes, and inflammation.¹⁻⁶ 36 37 The leaves of *C. paliurus* were approved as a new raw food material by The National 38 Health Commission of the People's Republic of China in 2013.⁷ Currently, C. paliurus is an important and valuable crop in the functional food market. A variety of 39 40 triterpenoid glycosides, flavonoids, and other compounds have been isolated from the leaves of this plant.⁸ Among its constituents, 3,4-seco-dammarane triterpenoids are 41 42 considered characteristic taxonomic indicators of this plant and the primary active and functional constituents in C. paliurus. To date, 20 seco-dammarane triterpenoid 43 glycosides, which are cleaved in plants by hydroxylation or methyl esterification at 44 C-3 and C-4 followed by glycosylation at C-12, have been reported.⁹⁻¹⁴ Some of them 45 have a sweet taste and are used as functional food ingredients. For example, 46 cyclocariosides I-II are approximately 50 to 100 times sweeter than 2% sucrose¹⁵⁻¹⁶ 47 and are used as sweeteners and antitumor agents.¹⁷ Although research on the 48 anti-inflammatory effects of an extract of C. paliurus has been reported,¹⁸ no active 49 anti-inflammatory compound has been reported. To explore potential bioactive 50

51 compounds from the leaves of C. paliurus, a bioactivity-guided investigation of anti-inflammatory compounds was carried out. The active fraction was isolated to 52 53 yield eleven new 3,4-seco-dammarane triterpenoid saponins (1–11) and seven known 54 dammarane triterpenoid saponins (12-18) (Figure 1). The inhibition of NO by the extracts and compounds were evaluated. Furthermore, the effects of the 55 anti-inflammatory compounds on the levels of inflammatory cytokines TNF- α , PGE₂ 56 57 and IL-6 and the expression of proteins iNOS, COX-2 and NF- κ B/p65 were tested using LPS-activated RAW 264.7 macrophage cells. We describe herein the separation 58 59 and structural identification of seco-dammarane triterpenoid saponins, their anti-inflammatory effects, and their structure-activity relationships. 60

61

62 MATERIALS AND METHODS

63 Instruments. The detailed experimental instruments are seen in the Support64 Information (Text S1).

Chemicals. Silica gel (Qingdao Marine Chemical Co., Ltd., China), Sephadex
LH-20 gel (Amersham Pharmacia Biotech AB, Sweden), and RP-C₁₈ (Merck,
Germany) were used for various separations. D-glucose, L-glucose, L-arabinose,
D-arabinose, D-quinovose, and L-quinovose were used as the sugar standards
(Sigma-Aldrich, Munich, Germany).

Plant materials. The leaves (20.0 kg) of *C. paliurus* were bought from Xiushui
county, Jiangxi Province, China in July 2017 and identified by Associate Professor
Qiang Xie (Guangxi Normal University). A specimen (No. ID-20170705) was
preserved in the State Key Laboratory of Medicinal Resources of Guangxi Normal
University.

75 **Preparation of extracts and isolation of triterpenoid saponins.** The fresh, dried

16 leaves (20.0 kg) of *C. paliurus* were extracted with 75% ethanol (EtOH, 4×3 h), and 177 the solvent was evaporated under vacuum to obtain an extract. The crude extract (2.0 178 kg) was suspended in distilled water and partitioned with polyethylene (PE), ethyl 179 acetate (EtOAc) and n-butanol (*n*-BuOH) to obtain a PE extract (480 g), an EtOAc 180 extract (450 g) and an *n*-BuOH extract (600 g). The anti-inflammatory effects were 181 analyzed by measuring the inhibition of NO production in LPS-activated RAW 264.7 182 cells.

83 The active EtOAc fraction was loaded to a chromatography column (CC) on silica 84 gel eluted with CH₂Cl₂-MeOH mixtures (from 1:0 to 0:1) to yield eight fractions (Fr. A to Fr. H). The anti-inflammatory fraction D (45 g) was fractionated using an MCI 85 gel CC eluting with MeOH-H₂O (from 3:7 to 1:0) to obtain Fr. D-1 to Fr. D-4. Fr. D-3 86 87 (10 g) was further separated using RP-C₁₈ CC eluting with gradients of MeOH-H₂O 88 (from 6:10 to 1:0) to afford Fr. D-3-1 to Fr. D-3-4. Fr. D-3-2 (0.8 g) was loaded to 89 Sephadex LH-20 eluting with gradients of H₂O-MeOH (from 1:0 to 0:1), and further 90 separated by RP-C₁₈ semi-preparative HPLC to furnish compounds 1 (28.0 mg), 2 91 (15.4 mg), and 3 (7.1 mg). Fr. D-3-3 (3.2 g) was applied to Sephadex LH-20 CC and 92 eluted with MeOH and then chromatographed with semi-preparative RP-C₁₈ HPLC to 93 afford compounds 4 (28.0 mg), 5 (15.2 mg), and 13 (30.3 mg). Fr. D-3-4 (4.2 g) was 94 chromatographed by silica gel and eluted with gradients of CH₂Cl₂-MeOH (from 1:0 95 to 0:1) to create Fr. D-3-4-1 to Fr. D-3-4-5. Fr. D-3-4-2 was subjected to Sephadex LH-20 CC and RP-C₁₈ HPLC to get compounds 6 (28.2 mg), 7 (15.2 mg), 11 (11.4 96 mg), and 12 (1.0 mg). In addition, Fr. D-3-4-3 was isolated by Sephadex LH-20 CC 97 98 and RP-C₁₈ HPLC to afford compounds 8 (28.2 mg), 9 (15.0 mg), and 10 (19.7 mg). 99 The anti-inflammatory Fr. E (80 g) was subjected to MCI gel eluting with gradients of 100 MeOH-H₂O (from 6:4 to 1:0) to yield Fr. E-1 to Fr. E-4. Active Fr. E-3 (23 g) was

101 further chromatographed by RP-C₁₈ CC eluted with gradients of MeOH-H₂O (from 6:10 to 1:0) to afford Fr. E-3-1 to Fr. E-3-4. Fr. E-3-2 was subjected to Sephadex 102 LH-20 CC and semi-preparative RP- C_{18} HPLC to afford compounds 14 (28.3 mg), 15 103 104 (15.4 mg), 16 (12.1 mg), 17 (12.0 mg), and 18 (12.3 mg). 105 *Cyclocarioside R (1)*: The various chromatographic isolation and purification gave 28.0 mg of white, amorphous powder; $[\alpha]_D^{25}$ + 38.0 (c 0.16, MeOH); negative 106 HRESIMS m/z calculated for C₃₆H₆₀ClO₁₀ [M + Cl]⁻, 687.3859, found: 687.3875; 107 ¹H- (500 MHz) and ¹³C-NMR (125 MHz) data (pyridine- d_5 , Tables 1 and 3). 108 109 Cyclocarioside S (2): The various chromatographic isolation and purification gave 15.4 mg of white, amorphous powder; $[\alpha]_D^{25}$ + 28.6 (c 0.12, MeOH); positive 110 HRESIMS m/z calculated for C₃₆H₆₀O₁₀Na [M + Na] +, 675.4073, found: 675.4084; 111 ¹H- (500 MHz) and ¹³C-NMR (125 MHz) data (pyridine- d_5 , Tables 1 and 3). 112 Cyclocarioside T(3): The various chromatographic isolation and purification gave 113 7.1 mg of white, amorphous powder; $[\alpha]_D^{25}$ + 10.6 (c 0.19, MeOH); negative 114 HRESIMS m/z calculated for $C_{37}H_{62}ClO_{10}$ [M + Cl]⁻, 701.4142, found: 701.4032; 115 ¹H- (500 MHz) and ¹³C-NMR (125 MHz) data (pyridine- d_5 , Tables 1 and 3). 116 117 *Cyclocarioside U (4)*: The various chromatographic isolation and purification gave 28.0 mg of white, amorphous powder; $[\alpha]_D^{25}$ + 27.6 (c 0.14, MeOH); positive 118

- 119 HRESIMS m/z calculated for C₃₆H₄₅O₄ [M + Na] ⁺, 641.4069, found: 641.4029; ¹H-
- 120 (500 MHz) and 13 C-NMR (125 MHz) data (pyridine- d_5 , Tables 1 and 3).
- 121 *Cyclocarioside V (5)*: The various chromatographic isolation and purification gave
- 122 15.2 mg of white, amorphous powder; $[\alpha]_D^{25}$ + 4.8 (c 0.14, MeOH); negative
- 123 HRESIMS m/z calculated for C₃₇H₆₀ClO₈ [M + Cl]⁻, 667.3960, found: 667.3977; ¹H-
- 124 (500 MHz) and 13 C-NMR (125 MHz) data (pyridine- d_5 , Tables 1 and 3).
- 125 *Cyclocarioside W* (6): The various chromatographic isolation and purification gave

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28.2 mg of white, amorphous powder; $[\alpha]_D^{25}$ + 11.3 (c 0.14, MeOH); positive HRESIMS m/z calculated for C₃₇H₆₀O₉Na [M + Na] +, 671.4136, found: 671.4135; 127 ¹H- (500 MHz) and ¹³C-NMR (125 MHz) data (pyridine- d_5 , Tables 1 and 3). 128 *Cyclocarioside X (7)*: The various chromatographic isolation and purification gave 129 15.2 mg of white, amorphous powder; $[\alpha]_D^{25}$ + 24.8 (c 0.07, MeOH); negative 130 HRESIMS m/z calculated for C₃₇H₆₂ClO₉ [M + Cl]⁻, 685.4074, found: 685.4082; ¹H-131 132 (500 MHz) and ¹³C-NMR (125 MHz) data (pyridine- d_5 , Tables 2 and 3). 133 Cyclocarioside Y (8): The various chromatographic isolation and purification gave 28.2 mg of white, amorphous powder; $[\alpha]_D^{25}$ + 1.5 (*c* 0.14, MeOH); positive 134 HRESIMS *m/z* calculated for C₃₈H₆₄O₉Na [M + Na] ⁺, 687.4491, found: 687.4448; 135 ¹H- (500 MHz) and ¹³C-NMR (125 MHz) data (pyridine- d_5 , Tables 2 and 3). 136 137 *Cyclocarioside* Z_1 (9): The various chromatographic isolation and purification gave 15.0 mg of white, amorphous powder; $[\alpha]_D^{25}$ + 2.9 (c 0.19, MeOH); positive 138 HRESIMS m/z calculated for C₃₇H₆₂O₉Na [M + Na] +, 673.4279, found: 673.4292; 139 ¹H- (500 MHz) and ¹³C-NMR (125 MHz) data (pyridine- d_5 , Tables 2 and 3). 140 Cyclocarioside Z_2 (10): The various chromatographic isolation and purification 141 gave 19.7 mg of white, amorphous powder; $[\alpha]_D^{25}$ + 34.0 (*c* 0.10, MeOH); negative 142 HRESIMS m/z calculated for C₃₇H₆₂ClO₉ [M + Cl]⁻, 685.4072, found: 685.4082; ¹H-143 144 (500 MHz) and ¹³C-NMR (125 MHz) data (pyridine- d_5 , Tables 2 and 3). 145 *Cyclocarioside* Z_3 (11): The various chromatographic isolation and purification gave 11.4 mg of white, amorphous powder; $[\alpha]_D^{25}$ + 38.9 (*c* 0.14, MeOH); negative 146 HRESIMS *m/z* calculated for C₃₈H₆₄ClO₉ [M + Cl]⁻, 699.422, found: 699.4239; ¹H-147 (500 MHz) and ¹³C-NMR (125 MHz) data (pyridine- d_5 , Tables 2 and 3). 148 149 Determination of the absolute configurations of the sugars. Each new compound was dissolved in 1 M HCl and stirred at 80 °C for 2 hours. The solution of 150

the reactant was then extracted with EtOAc 3 times. The H₂O layers were evaporated under vacuum. The presence of L-arabinose in the aqueous layers of compounds **1**, **2**, **4**, **7**, and **10** were detected by Sigel TLC (H₂O-MeOH-CH₂Cl₂, 1:8:12) with standard sample. D-quinovose was confirmed as the sugar moiety of compounds **3**, **5**, **8**, **9** and **15 11** by Sigel TLC (H₂O-MeOH-CH₂Cl₂, 1:8:12) with standard sample, and D-glucose was identified as the sugar moiety of compound **6** by Sigel TLC (H₂O-MeOH-CH₂Cl₂, 1:8:12) with standard sample.

Cell viability. RAW 264.7 cell viability was performed using the previously
described MTT method.¹⁹ The detailed procedures are seen in the Support
Information (Text S2).

NO production assay. NO production was detected according to a previous
 report.²⁰ The detailed procedures are seen in the Support Information (Text S3).

163 Enzyme-linked immunosorbent assay (ELISA). The expression of 164 proinflammatory cytokines (TNF- α , PGE₂, and IL-6) was tested using the previously 165 described method.²¹ The detailed procedures are seen in the Support Information 166 (Text S4).

167 Western blotting analysis. The expressions of proteins iNOS, NF- κ B/p65, and 168 COX-2 were assayed by Western blotting analysis according to standard procedures 169 published previously.¹⁹ The detailed procedures are seen in the Support Information 170 (Text S5).

171 Statistical analysis. All data were performed by Student's test. The results are 172 displayed as the mean \pm standard deviation (SD) for three independent experiments. 173 Differences were considered statistically significant at p < 0.05.

174

175 **RESULTS AND DISCUSSION**

176 Structural elucidation of the 3,4-*seco*-dammarane triterpenoid saponins.

Cyclocarioside R (1) yielded a negative-ion peak $[M + Cl]^{-}$ at m/z 687.3859 (calcd. 177 for C₃₆H₆₀ClO₁₀, 687.3875) in its HRESIMS spectrum (Figure S1), which 178 demonstrated a molecular formula, $C_{36}H_{60}O_{10}$, in light of its 1D NMR (pyridine- d_5) 179 and HRESIMS data. The ¹H NMR data (pyridine- d_5 , Table 1) of 1 exhibited six 180 181 methyl signals at $\delta_{\rm H}$ 0.96 (s, 30-CH₃), 1.02 (s, 18-CH₃), 1.24 (s, 19-CH₃), 1.58 (s, 182 21-CH₃), 2.06 (s, 27-CH₃), and 1.80 (s, 29-CH₃); a methoxy group at $\delta_{\rm H}$ 3.51 (s, 3-OMe); two terminal methylene olefinic protons at $\delta_{\rm H}$ 5.33 (br s, H-26a), 5.15 (br s, 183 184 H-26b), 4.87 (br s, H-28a), and 4.95 (br s, H-28b); and a glycosyl anomeric proton at $\delta_{\rm H}$ 4.80 (d, $J_{\rm H-1'/H-2'}$ = 7.4 Hz, H-1'). The ¹³C-NMR (pyridine-d₅, Table 3) and the 185 distortionless enhancement by polarization transfer (DEPT) (Figure S4) spectra 186 displayed the existence of a carboxyl group (-C=O) at $\delta_{\rm C}$ 176.3 and two terminal 187 188 double bonds (-C=CH₂) at $\delta_{\rm C}$ 148.9 (C-4), 114.3 (C-28) and $\delta_{\rm C}$ 147.8 (C-25), 113.5 (C-26). The ¹H-¹H correlation spectroscopy (COSY) (Figures 2 and S6) of 1 189 190 demonstrated the presence of four fragments (H-1/H-2, H-5/H-6/H-7, H-11/H-12/ 191 H-13/H-17/H-16/H-15, H-22/H-23/H-24). In its HMBC spectrum (Figures 2 and S7), a key correlation from H-3-OMe ($\delta_{\rm H}$ 3.51, s) to C-3 ($\delta_{\rm C}$ 176.3) suggested that the – 192 OCH₃ connected to C-3. The cross peaks from H-28 ($\delta_{\rm H}$ 4.95, 4.87, br s) to C-5 ($\delta_{\rm C}$ 193 52.5) and C-29 ($\delta_{\rm C}$ 24.2, s) and from H-26 ($\delta_{\rm H}$ 5.33, 5.15, br s) to C-24 ($\delta_{\rm C}$ 80.9) and 194 195 C-27 ($\delta_{\rm C}$ 19.1) indicated the two terminal double bonds (–C=CH₂) linked to C-4 and C-25, respectively. The HMBC correlations from H-21-CH₃ ($\delta_{\rm H}$ 1.58, s) to C-17 ($\delta_{\rm C}$ 196 52.3) confirmed the side chain linked to C-17. The HMBC correlations from the H-1' 197 198 ($\delta_{\rm H}$ 4.80, d, anomeric proton) to C-12 ($\delta_{\rm C}$ 75.7) suggested that the arabinopyranose was linked to C-12. Its diagnostic ROESY correlations of H-5/H-9/H-12/H-17/ 199 200 H-CH₃-21/H-23/H-CH₃-30 suggested that H-5, H-9, H-12, H-17, CH₃-21, H-23, and 201 CH₃-30 are α -oriented (Figures 3 and S8). The β -orientations of H-13, H-CH₃-18, and H-CH₃-19 were then verified by the ROESY correlations of H-13/H-CH₃-18 202 203 /H-CH₃-19. In addition, the pentose unit was assigned as α -L-arabinopyranosyl based 204 on its ¹H-and ¹³C-NMR data (pyridine- d_5 , Tables 1 and 3) and comparison with other 3,4-seco-dammarane triterpenoid saponin from this plant¹¹⁻¹³ as well as further 205 206 hydrolysis of 1 with 1M HCl and Sigel TLC with standard L-arabinose. The coupling 207 constant of $J_{\text{H-1'/H-2'}} = 7.4$ Hz of the anomeric proton suggested that the L-arabinose was also α -oriented. Analyses of the 1D and 2D-NMR data of 1 showed high 208 similarity to those of cyclocarioside K,¹⁴ suggesting that 1 includes the basic 209 3,4-seco-dammarane triterpenoid skeleton, differing from cyclocarioside K only in the 210 signals of the side chain. The 20-, 23- and 24-OH substituents were confirmed by the 211 212 ¹H-¹H COSY correlations of H-22/H-23/H-24 along with the key HMBC correlations 213 from H-21-CH₃ to C-20 ($\delta_{\rm C}$ 76.2) and C-22 ($\delta_{\rm C}$ 41.9) and from H-23 to C-20 ($\delta_{\rm C}$ 76.2), C-24 ($\delta_{\rm C}$ 80.9) and C-25 ($\delta_{\rm C}$ 147.8) (Figure 2). The above information indicated 214 215 that 1 was a characteristic 3,4-seco-dammarane saponin.

The molecular formula of cyclocarioside S (2) was deduced to be $C_{36}H_{60}O_{10}$ based 216 on its sodium adduct molecular ion peak $[M + Na]^+$ at m/z 675.4073 (calcd. for 217 218 $C_{36}H_{60}O_{10}Na$, 675.4084) in its HRESIMS (Figure S9) spectrum, which has the same 219 formula as that of 1. Compound 2 had almost identical NMR (Tables 1 and 3) data to 220 that of 1, as revealed by comparisons of their MS and NMR signals. Small differences 221 existed in the ¹H- and ¹³C-NMR (pyridine- d_5) data of positions 22, 23, and 24 between 2 and 1. The ¹H NMR signals of H-22, 23, and 24 shifted from $\delta_{\rm H}$ 1.90, 2.17, 222 223 4.54, and 4.40 in 1 to $\delta_{\rm H}$ 2.12, 2.29, 4.61, and 4.57 in 2, respectively (Table 1). Moreover, the ¹³C NMR signals of C-22, 23, and 24 shifted from $\delta_{\rm C}$ 41.9, 71.8, and 224 225 80.9 in 1 to $\delta_{\rm C}$ 40.1, 71.6, and 79.9 in 2, respectively (Table 3). Furthermore, compounds 1 and 2 exhibited two peaks with different retention times in the HPLC
chromatogram (Figure S17), indicating that 1 and 2 are epimers. The results indicate
that 2 is also a 3,4-*seco*-dammarane triterpenoid saponin.

229 To assess the relative configurations of positions 23 and 24 in their side chains of 1 and 2, their ROESY correlations and ¹H-¹H coupling constant values were focused. 230 The ROESY correlation of H-23/H-CH₃-21 indicated that H-23 and H-CH₃-21 are 231 232 α -oriented (Figures 3 and S8). Moreover, the ROESY spectrum of 2 (Figures 3 and S16) showed a correlation between H-24/H-CH₃-21, indicating that H-CH₃-21 and 233 H-24 are α -oriented. Furthermore, piscidinol A²² and 24-*epi*-piscidinol A,²³ which 234 235 possessed the same eight-carbon side chains as 1 and 2 with syn and anti orietations 236 between hydrogens at positions 23 and 24, revealed coupling constants of $J \approx 0$ Hz 237 and 8.0 Hz, and their structures were identified by X-ray analysis. Correspondingly, combretanone A,²⁴ alisol E²⁵ and 23,24,25-trihydroxycycloartan-3-one,²⁶ which have 238 syn and anti orietations between hydrogens at positions 23 and 24, exhibited coupling 239 240 constants of $J \approx 0$ Hz and 6.0 Hz, respectively. The similar coupling constants of respective J = 6.4 Hz and J = 5.8 Hz of 1 and 2 between hydrogens at positions 23 241 and 24 suggested that both hydrogens linked to positions 23 and 24 were in 242 anti-configurations. Thus, on the basis of their ROESY correlations and coupling 243 244 constants between hydrogens at positions 23 and 24, the absolute configurations of 245 carbons at positions 23 and 24 of 1 and 2 were determined as 23R, 24S, and 23S, 24R, 246 respectively. Compounds 1 and 2 were named cyclocarioside R and cyclocarioside S, 247 respectively (Figure 1).

248 Cyclocarioside T (**3**) yielded a negative-ion peak $[M + Cl]^-$ at m/z 701.4142 (calcd. 249 for C₃₇H₆₂ClO₁₀, 701.4132) in its HRESIMS spectrum (Figure S18), which 250 demonstrated a molecular formula, C₃₇H₆₂O₁₀, in light of its 1D NMR (pyridine- d_5) 251 and HRESIMS data. The ¹H-, ¹³C-NMR data (pyridine- d_5 , Tables 1 and 3) of **3** were similar to those of compound 1, indicating that 3 also possesses a similar 252 3,4-seco-dammarane triterpenoid skeleton as 1, differing only in the presence of a 253 254 β -D-quinovopyranosyl unit at position 12 and the absence of an α -L-arabinopyranosyl unit. The sugar unit was identified as a β -D-quinovopyranosyl moiety based on the 255 ¹H-, ¹³C-NMR data (pyridine- d_5 , Tables 1 and 3), the hydrolysis of **3** with 1M HCl 256 257 and Sigel TLC with standard sample. In its HMBC spectrum (Figures 2 and S24), the methyl singlets H-6'-CH₃ ($\delta_{\rm H}$ 1.59, d, J = 5.6 Hz) correlated with an oxymethine 258 259 carbon resonance at C-5' ($\delta_{\rm C}$ 73.3), indicating that H-6'-CH₃ was connected to C-5', and H-1' ($\delta_{\rm H}$ 4.88, d, $J_{\rm H-1'/H-2'}$ = 7.9 Hz, a glycosyl anomeric proton) was correlated 260 with C-12 ($\delta_{\rm C}$ 76.3, an oxymethine carbon resonance), suggesting the sugar group was 261 262 connected to C-12. In the ROESY spectrum of 1, the correlation of H-CH₃-21/H-23 263 indicated that H-CH₃-21 and H-23 are α -oriented (Figures 3 and S25). Moreover, the coupling constant of J = 6.2 Hz of **3** suggested that H-23 and H-24 were in 264 265 anti-configurations with respect to their configurations in 1, and the absolute 266 configurations of carbons at positions 23 and 24 of **3** was assigned as 23R, 24S. Thus, **3** was determined and named cyclocarioside T (Figure 1). 267

Cyclocarioside U (4) yielded a positive-ion peak $[M + Na]^+$ at m/z 641.4069 268 269 (calcd. for C₃₆H₅₈O₈Na, 641.4029) in its HRESIMS spectrum (Figure S26), which 270 demonstrated a molecular formula, $C_{36}H_{58}O_8$, in light of its 1D NMR (pyridine- d_5) 271 and HRESIMS data. Its ¹H-NMR data (pyridine- d_5 , Table 1) exhibited six methyl 272 signals at $\delta_{\rm H}$ 0.95 (s, 30-CH₃), 1.05 (s, 18-CH₃), 1.25 (s, 19-CH₃), 1.49 (s, 21-CH₃), 273 1.89 (s, 27-CH₃), and 0.95 (s, 29-CH₃); a methoxy signal at $\delta_{\rm H}$ 3.52 (s, 3-OMe); three olefinic protons including two terminal double bond protons at $\delta_{\rm H}$ 6.15 (m, H-23), 274 275 6.45 (d, J = 15.7 Hz, H-24); $\delta_{\rm H}$ 5.07 (br s, H-26a), 4.98 (br s, H-26b); and $\delta_{\rm H}$ 4.87 (br

s, H-28a), 4.95 (br s, H-28b); and a glycosyl anomeric proton at $\delta_{\rm H}$ 4.82 (d, $J_{\rm H-1'/H-2'}$ = 276 7.4 Hz, H-1'). Its spectra of ¹³C-NMR (pyridine- d_5 , Table 3) and DEPT (Figure S29) 277 indicated the presence of a carboxyl unit at $\delta_{\rm C}$ 176.3 (C-3) and three double bond 278 279 signals at $\delta_{\rm C}$ 148.9 (C-4), 114.2 (C-28); $\delta_{\rm C}$ 128.7 (C-23), 135.9 (C-24); and $\delta_{\rm C}$ 143.1 (C-25), 115.4 (C-26). The ¹H-, ¹³C-NMR data (pyridine- d_5) for 4 displayed a number 280 281 of similarities to those of 1 but differed in the signals of the eight-carbon side chain, 282 indicating that 4 also possesses a 3,4-seco-dammarane triterpenoid skeleton. Its ¹H-¹H COSY (Figures 2 and S31) correlations of H-22 ($\delta_{\rm H}$ 2.58, 2.71, m)/H-23 ($\delta_{\rm H}$ 6.15, 283 284 m)/H-24 ($\delta_{\rm H}$ 6.45, d, J = 15.7 Hz) suggested the existence of a -CH₂CH=CHfragment. The HMBC (Figures 2 and S32) correlations of H-23 ($\delta_{\rm H}$ 6.15, m) with 285 C-20 ($\delta_{\rm C}$ 74.9) and H-24 ($\delta_{\rm H}$ 6.45, d, J = 15.7 Hz) with C-26 ($\delta_{\rm C}$ 115.4) and C-27 ($\delta_{\rm C}$ 286 287 19.3) indicated the location of the additional double bond is at positions 23 and 24. In 288 addition, the sugar unit was identified as α -L-arabinopyranosyl based on the ¹H-, ¹³C-NMR (pyridine- d_5 , Tables 1 and 3) data, the hydrolysis of 4 with 1M HCl and 289 290 Sigel TLC with L-arabinose, as well as comparison with those of 1. The L-arabinose 291 is linked to C-12 by the HMBC correlation from H-1' (anomeric proton) to C-12. Therefore, the structure of 4 was identified and assigned as cyclocarioside U (Figure 292 293 1).

The molecular formula of cyclocarioside V (**5**), $C_{37}H_{60}O_8$, was determined by 1D NMR (pyridine- d_5) data and HRESIMS m/z 667.3960 [M + Cl] ⁻ (calcd. for $C_{37}H_{60}ClO_8$, 667.3977) (Figure S34), which is 14 Da more than that of **4**. The 1D NMR (pyridine- d_5 , Tables 1 and 3) signals of **5** closely resembled those of **4**; they differed in the existence of a –CH₃ at δ_H 1.55 (d, J = 5.6 Hz). Comparisons of the 1D NMR, MS data between **5** and **4** revealed a pronounced difference in the signals for the sugar moiety and the presence of a methyl hexacarbonose in **5**, indicating that **5** 301 also possesses a 3,4-seco-dammarane triterpenoid skeleton. The hydrolysis of 5 with 1M HCl and Sigel TLC with standard sample attributed the sugar in 5 as a 302 303 quinovopyranose. The arabinopyranose linked to position 12 in 4 was replaced by a quinovopyranose in 5. The coupling constant of $J_{H-1'/H-2'} = 7.3$ Hz of the anomeric 304 305 proton, suggested the quinovopyranose was β -oriented. In the HMBC spectrum, the 306 correlation of H-6'-CH₃ ($\delta_{\rm H}$ 1.55, d, J = 5.6 Hz) with C-5' ($\delta_{\rm C}$ 73.2) indicated 6'-CH₃ connected to C-5'. The HMBC correlation (Figures 2 and S40) of H-1' (anomeric 307 308 proton) with C-12 unambiguously confirmed that the sugar unit was situated at C-12. Thus, the structure of 5 was deduced and given the common name cyclocarioside V 309 (Figure 1). 310

311 Cyclocarioside W (6) yielded a negative-ion peak $[M + Cl]^{-1}$ at m/z 671.4136 (calcd. for C₃₇H₆₀ClO₁₀, 671.4135) in its HRESIMS spectrum (Figure S42), which 312 demonstrated a molecular formula, $C_{37}H_{60}O_{10}$, in light of its 1D NMR (pyridine- d_5) 313 and HRESIMS data. The 1D NMR (pyridine- d_5 , Tables 1 and 3) spectra of 6 were 314 closely similar to those of 5 except for the absence of the –CH₃ peaks [$\delta_{\rm H}$ 1.55 (3H, d, 315 316 5.9 Hz); $\delta_{\rm C}$ 18.9] observed in 5 and the existence of alternative –CH₂OH singlets [$\delta_{\rm H}$ 317 4.38, 4.52 (2H, m); $\delta_{\rm C}$ 63.9] in 6, suggesting that 6 is also a 3,4-seco-dammarane 318 triterpenoid. In the HMBC spectrum (Figures 2 and S48), the correlation of 6'-CH₂OH 319 $(\delta_{\rm H} 4.38, 4.52, \text{ m})$ with C-5' $(\delta_{\rm C} 72.8)$ indicated that the -CH₂OH was linked to C-5'. The HMBC correlation of H-1' ($\delta_{\rm H}$ 4.94, anomeric proton) with C-12 ($\delta_{\rm C}$ 76.6) 320 indicated the sugar was connected to C-12. The hydrolysis of 6 with HCl yielded 321 D-glucose as identified by Sigel TLC analysis. The coupling constant of $J_{\text{H-1'/H-2'}} = 7.3$ 322 Hz of the glucosyl anomeric proton indicated the β -orientation for 6. Consequently, 323 the structure of 6 was deduced and given the common name cyclocarioside W (Figure 324 325 1).

326 Cyclocarioside X (7) yielded a positive-ion peak $[M + Na]^+$ at m/z 673.427 in its HRESIMS spectrum (Figure S50), which demonstrated a molecular formula, 327 328 $C_{37}H_{62}O_9$, in light of its 1D NMR (pyridine- d_5) and HRESIMS data. The 1D NMR 329 spectra (pyridine- d_5 , Tables 2 and 3) of 7 were closely analogous to those of 4 except 330 in the signals of their side chains, indicating that 7 also possesses a 3,4-seco-dammarane triterpenoid skeleton. There were signals from a methyl group 331 332 $(\delta_{\rm H} 1.34, s; \delta_{\rm C} 26.4, 26$ -CH₃), a quaternary carbon $(\delta_{\rm C} 74.6, C$ -25) and an additional methoxy group ($\delta_{\rm H}$ 3.24, s; $\delta_{\rm C}$ 50.7, 25-OCH₃) in 7, replacing the terminal olefin at $\delta_{\rm H}$ 333 334 4.95 and 5.01, and at $\delta_{\rm C}$ 115.4 and 143.1 observed in 4. The HMBC (Figures 2 and S56) correlations from $\delta_{\rm H}$ 3.24 (25-OCH₃) and $\delta_{\rm H}$ 1.34 (CH₃-26/27) to $\delta_{\rm C}$ 74.6 (C-25) 335 confirmed that the methoxy group and gem-dimethyl were linked to position C-25. 336 337 The hydrolysis of 7 with 1M HCl yielded L-arabinose. The coupling constant of $J_{\text{H-1'/H-2'}} = 7.4$ Hz of the sugar anomeric proton indicated the α -orientation for the 338 arabinosyl group. Accordingly, the structure of 7 was established and given the 339 340 common name cyclocarioside X (Figure 1).

341 Cyclocarioside Y (8) was revealed to have the molecular formula $C_{38}H_{64}O_9$ by the NMR data (¹H, ¹³C, pyridine- d_5) and its positive-ion peak [M + Na] ⁺ at m/z 687.4491 342 (calcd. for C₃₈H₆₄O₉Na, 687.4448) (Figure S58) in the HRESIMS spectrum, which is 343 344 14 Da more than that of 7. The NMR data (¹H, ¹³C, pyridine- d_5 , Tables 2 and 3) of 8 345 were closely analogous to those of 7; they differed in the existence of a methyl group signals H-6'-CH₃ ($\delta_{\rm H}$ 1.59, d, 6.0 Hz; $\delta_{\rm C}$ 18.9) in 8, indicating that 8 is also a 346 3,4-seco-dammarane triterpenoid saponin. In the HMBC spectrum, the correlation of 347 348 H-6'-CH₃ ($\delta_{\rm H}$ 1.59, d, J = 6.0 Hz) with C-5' ($\delta_{\rm C}$ 73.2) indicated that H-6'-CH₃ is linked to C-5'. The HMBC correlation (Figures 2 and S40) of H-1' ($\delta_{\rm H}$ 4.94) with C-12 ($\delta_{\rm C}$ 349 350 76.3) indicated the linkage of the sugar unit to the 3,4-seco-dammarane triterpenoid

351 skeleton was at C-12. The hydrolysis with 1M HCl yielded quinovopyranose. The 352 coupling constant of $J_{\text{H-1'/H-2'}} = 7.2$ Hz suggested the quinovopyranose was β -oriented. 353 Thus, the structure of **8** was elucidated and given a trivial name cyclocarioside Y 354 (Figure 1).

Cyclocarioside Z_1 (9) was found to have the formula $C_{37}H_{62}O_9$ via the NMR data 355 (¹H, ¹³C, pyridine- d_5) and its negative-ion peak [M + Cl] ⁻ at m/z 685.4074 (calcd. for 356 357 $C_{37}H_{62}ClO_{9}$, 685.4082) (Figure S66) in the HRESIMS, which is 14 Da less than that of 8. The NMR data (¹H, ¹³C, pyridine- d_5 , Tables 2 and 3) of 9 showed that the signals 358 359 were analogous to those of 8 except for the absence of an –OCH₃ signals ($\delta_{\rm H}$ 3.23/ $\delta_{\rm C}$ 50.7) in 9, suggesting that 9 is also a 3,4-seco-dammarane triterpenoid saponin. The 360 NMR data (¹H, ¹³C, pyridine- d_5 , Tables 2 and 3, Figures 2 and S72) confirmed that an 361 362 $-OCH_3$ at position C-25 in 9 was substituted with a -OH in 8. The sugar unit was confirmed by hydrolysis to be quinovopyranose. The coupling constant of $J_{H-1/H-2'}$ = 363 7.2 Hz demonstrated that the quinovopyranose was β -oriented. Thus, the structure of 9 364 365 was identified and given the common name cyclocarioside Z_1 (Figure 1).

Cyclocarioside Z_2 (10) has the same molecular formula of $C_{37}H_{62}O_9$ as 7, as 366 evidenced by the NMR (¹H, ¹³C, pyridine- d_5) data and its negative-ion peak [M + Cl] 367 ⁻ at m/z 685.4072 (calcd. for C₃₇H₆₂ClO₉, 685.4082) (Figure S74) in the HRESIMS 368 369 spectrum. The ¹H- and ¹³C-NMR data (pyridine- d_5 , Tables 2 and 3) of 10 were 370 analogous to those of 7 except for the positions of the double bond and the –OH of the 371 side chain, indicating that 10 also possesses a 3,4-seco-dammarane skeleton. In its ¹H-¹H COSY spectrum (Figures 2 and S79), the correlations of H-22/H-23/H-24 372 373 suggested that there is a -CH₂CH(O)CH=C fragment. In its HMBC spectrum (Figures 2 and S80), the correlations of H-23-OCH₃ ($\delta_{\rm H}$ 3.24, s) with C-23 ($\delta_{\rm C}$ 76.3) and H-24 374 375 $(\delta_{\rm H} 5.22, d, J = 6.0 \text{ Hz})$ with C-25 $(\delta_{\rm C} 135.9)/\text{C-26} (\delta_{\rm C} 18.7)/\text{C-27} (\delta_{\rm C} 26.2)$ indicated that the methoxy group and the double bond were located at C-23 and C-24/C-25, respectively. The hydrolysis of **10** yielded L-arabinose. The α -configuration for the sugar was confirmed by its coupling constant of 7.4 Hz between H-1' and H-2'. These findings indicated that **10** possessed a skeleton of 3,4-*seco*-dammarane triterpenoid.

Cyclocarioside Z_3 (11) has the molecular formula $C_{38}H_{64}O_9$ (Figure S82) by the 380 NMR (¹H, ¹³C, pyridine- d_5) data and its negative-ion peak [M + Cl] - at m/z 699.4222 381 382 (calcd. for $C_{38}H_{64}ClO_{9}$, 699.4239) in the HRESIMS spectrum, which is 14 Da more than that of 10. The NMR (¹H, ¹³C, pyridine- d_5 , Tables 2 and 3) spectra of 11 383 384 suggested that its basic signals were analogous to those of 10 except for the existence of a β -D-quinovopyranosyl unit at C-12 and the absence of an α -L-arabinopyranosyl 385 moiety at C-12. The sugar unit was determined to be a β -D-quinovopyranosyl moiety 386 387 based on the hydrolysis of 11 with 1M HCl and Sigel TLC with D-quinovose and the 388 NMR (¹H, ¹³C, pyridine-d₅, Tables 2 and 3) data. The HMBC (Figures 2 and S88) correlation of H-6'-CH₃ ($\delta_{\rm H}$ 1.58) with C-5' ($\delta_{\rm C}$ 73.2) suggested H-6'-CH₃ linked to 389 390 C-5'. The HMBC correlation of H-1' ($\delta_{\rm H}$ 4.94, anomeric proton) with C-12 ($\delta_{\rm C}$ 76.2) indicated that the linkage of the sugar unit to the 3,4-seco-dammarane skeleton was at 391 C-12. The results revealed that 11 was also a 3,4-seco-dammarane triterpenoid 392 saponin. 393

394 To determine the relative configurations of position 23 of 10 and 11, their ROESY 395 spectra and comparisons of the NMR (¹H, ¹³C, pyridine- d_5) data with two known 396 epimers which possess the same side chain were focused. In the ROESY spectra of 10 and 11 (Figures 3, S81 and S89), the correlations of H-23/H-21-CH₃ indicated that 397 H-23 in each compound was α -oriented. In addition, the ¹³C-NMR data at $\delta_{\rm C}$ 127.4 398 (C-24), $\delta_{\rm C}$ 135.9 (C-25) of **10** and at $\delta_{\rm C}$ 127.5 (C-24), $\delta_{\rm C}$ 136.4 (C-25) of **11** more 399 400 closely resembled those 128.4 (C-24), $\delta_{\rm C}$ 135.6 (C-25) at δ_{C} of 401 23β -cycloarta-24-ene- 3β ,23-diol than those at $\delta_{\rm C}$ 129.1 (C-24), $\delta_{\rm C}$ 133.8 (C-25) of 402 23α -cycloarta-24-ene- 3β ,23-diol.²⁷ From the above analyses, the methoxy groups in 403 position 23 of the side chains of **10** and **11** were deduced to be β -oriented. 404 Correspondingly, the absolute configurations of position 23 in **10** and **11** were 405 unambiguously confirmed to be *R*. Compounds **10** and **11** were designated 406 cyclocarioside Z₂ and cyclocarioside Z₃, respectively (Figure 1).

By comparisons of the NMR data (¹H, ¹³C) to those reported in the literatures, the known isolates were identified as cyclocariol H (**12**),¹⁰ cyclocarioside O (**13**),²⁸ cyclocarioside P (**14**),²⁸ cyclocarioside J (**15**),²⁹ cyclocarioside I (**16**),¹⁶ cyclocarioside H (**17**),¹¹ and cyclocarioside K (**18**).¹²

411 The extracts and the compounds suppress the release of NO in LPS-activated RAW 264.7 cells. As shown in Figure 4, relative to the PE, EtOAc, n-Butanol, and 412 H₂O extracts, the EtOAc fraction of the 75% EtOH-H₂O extract revealed strong 413 inhibition to the release of NO in LPS-activated RAW 264.7 cells, comparable to the 414 achieved by the positive control dexamethasone. 415 inhibition Thus. the 416 anti-inflammatory compounds were further isolated from the EtOAc fraction.

The cell viability assay showed that these compounds did not exhibit any cytotoxicity, and the survival rates of RAW 264.7 cells were over 90% at a concentration of 40 μ M of each compound. Studies have shown after RAW 264.7 cells are treated with LPS, large amounts of NO, TNF- α , PGE₂ and IL-6 can be released, which result in inflammation.³⁰⁻³²

The inhibitory activities of 1-18 against LPS-activated NO release in RAW 264.7 cells are shown in Table 4. From the results, it was deduced that compounds 2, 4, 6-8, 10-13, and 16-17 suppressed the release of NO, with IC₅₀ values ranging from 8.23 μ M to 34.57 μ M. Compounds 7, 8, 10, and 11 showed strong inhibition to the 426 production of NO with IC₅₀ values of 8.23 μ M, 9.41 μ M, 9.36 μ M, and 11.23 μ M, 427 respectively, comparable to the inhibition achieved by the positive control 428 dexamethasone (IC₅₀: 8.34 μ M). Compounds 1, 3, 5, 9, 14-15, and 18 showed weak 429 inhibition of the release of NO, with IC₅₀ values above 40 μ M.

430 A comparison of the structures of these dammarane triterpenoid saponins with their activities revealed that the methoxy group in the side chain seemed to be related to 431 432 high potency against LPS-activated NO release in RAW 264.7 cells. For example, among the compounds, compounds 7, 8, 10, and 11, with methoxy groups at positions 433 434 25 or 23, achieved the most effective inhibition of NO release. Moreover, the dammarane triterpenoid saponins (1-6, 9, and 12-18), without methoxy groups on the 435 436 side chain, achieved reduced or no inhibition of the release of NO. Furthermore, by 437 comparing the inhibitory activities against NO release with their structures, it was 438 deduced that the kind of sugar group was important in influencing the inhibitory effect on NO release. Compounds 2, 4, 7, and 10, with an arabinopyranosyl moiety at 439 440 position 12, exhibited stronger activity against the release of NO than their corresponding compounds 3, 5, 8, and 11, with a quinovopyranosyl moiety at position 441 442 12. Similarly, compounds 13, with two arabinopyranosyl moieties at positions 3 and 12, revealed stronger inhibitory activities against the release of NO than their 443 444 corresponding compounds 14, with an arabinopyranosyl unit and a quinovopyranosyl 445 unit at positions 3 and 12, respectively. Furthermore, the inhibitory activities against 446 the release of NO were related to the absolute configurations of the side chain at positions 23 and 24. For example, compound 2 possesses different absolute 447 448 configurations of the side chain at positions 23 and 24 than compounds 1 and 3 and showed higher activity than both 1 and 3. The inhibitory activity against the release of 449 450 NO was also found to be related to the substitution positions of the double bond and -

451 OCH₃ in the side chain. For example, the substitution positions of the double bond and -OCH₃ in the side chains of 7 and 8 differ from those of 10 and 11 and showed 452 different activity. Comparison of the structures of 4 and 13 revealed that the inhibitory 453 activity against NO release appears largely unrelated to the presence of cleavage in 454 positions 2 and 3. In addition, comparison of the structures of 13 and 16 indicated that 455 the inhibitory activity is related to the existence of a ring in the side chain. For 456 457 example, compound 13, without an oxygen-substituted five-member ring in the side chain, revealed weaker inhibitory effects on NO release than did compound 16, which 458 459 has a ring. In summary, the inhibitory effects of dammarane triterpenoid saponins 1-18 on the release of NO depend on the substitution of the –OCH₃ on the side chain, 460 the kind of sugar, the absolute configuration of the side chain, the substitution 461 462 positions of the double bond and -OCH₃, the cleavage in positions 2 and 3, and the 463 presence of a ring in the side chain.

Active compounds reduce the expression of $TNF-\alpha$, PGE_2 and IL-6 in 464 465 LPS-activated RAW 264.7 cells. To explore the inhibition to inflammation of active compounds 7, 8, 10, and 11, three crucial proinflammatory mediators, TNF- α , PGE₂ 466 and IL-6, were investigated whether these compounds are able to decrease these 467 proinflammatory molecules stimulated by LPS. The levels of cytokines TNF- α , PGE₂, 468 469 and IL-6 markedly increased after LPS treatment. However, the overexpression of 470 cytokines TNF- α , PGE₂, and IL-6 was reduced by treatment with 7, 8, 10, and 11 471 (Figures 5, 6, and 7). Generally, the results suggested that these isolates can inhibit the expression of cytokines TNF- α , PGE₂, and IL-6 in a concentration-dependent manner 472 473 in LPS-activated RAW 264.7 cells. Notably, among the inhibitory effects of the four 474 isolates on the expression levels of cytokines TNF- α , PGE₂, and IL-6, those of 475 compound 7 were the most significant. The inhibitory activity of 7 was almost 476 equivalent to that of the positive control, dexamethasone.

Compound 7 inhibits the levels of proteins COX-2, iNOS, and NF- κ B/p65 in 477 LPS-activated RAW 264.7 cells. NF- κ B is a critical transcription factor regulating 478 proinflammatory cytokine expression during inflammation.³³⁻³⁶ Moreover, proteins 479 iNOS and COX-2 are important proinflammatory mediators that contribute to 480 inflammation.³⁷⁻⁴⁰ To detect the effects of active compound 7 on protein and gene 481 expression, the levels of COX-2, iNOS, and NF- κ B/p65 in LPS-activated cells were 482 measured. Activated by LPS, RAW 264.7 cells exhibited markedly increased levels of 483 484 the proteins COX-2, iNOS, and NF- κ B/p65 relative to the levels of control cells (Figure 8). After treatment with compound 7, the levels of proteins COX-2, iNOS, 485 and NF-*k*B/p65 in LPS-activated cells were markedly decreased. Compound 7 486 487 inhibited the expression levels of proteins COX-2, iNOS, and NF- κ B/p65 in 488 LPS-activated RAW 264.7 cells in a dose-dependent manner.

Conclusion. The C. paliurus is a famous edible medicinal plant in China, and its 489 490 potent anti-inflammatory effects have attracted increasing attention. However, before 491 the present study, its active anti-inflammatory compounds had not been identified. In 492 the present study, anti-inflammatory-guided isolation of the leaves of C. paliurus identified eighteen dammarane triterpenoid glycosides, including eleven new 493 494 3,4-seco-dammarane triterpenoid glycosides (1-11). Bioassays demonstrated that 495 compounds 7-8 and 10-11 showed potent inhibitory effects on the expression of cytokines NO, TNF- α , PGE₂, and IL-6. In addition, compound 7 significantly 496 suppressed the expression levels of proteins COX-2, iNOS, and NF-kB/p65, 497 498 indicating that this compound produced anti-inflammatory effects through the suppression of NF- κ B signaling pathways. The structure-activity relationship analysis 499 500 suggested that the anti-inflammatory activities of dammarane triterpenoid saponins are

501 correlated with the substitution of the methoxy group in the side chain, the kind of sugar, and the absolute configuration of the side chain of the compounds. In summary, 502 C. paliurus contains abundant anti-inflammatory dammarane triterpenoid saponins, 503 504 and compound 7 is a promising natural anti-inflammatory drug that warrants further investigation. These findings suggest that regular consumption of the leaves of C. 505 506 paliurus may help prevent the occurrence of inflammatory diseases. 507 **ASSOCIATED CONTENT** 508 509 **Supporting Information** The Supporting Information is available free of charge on the ACS Publications 510 website at DOI: xxxxxx. ¹H NMR, ¹³C NMR, DEPT, HSQC, ¹H-¹H COSY, HMBC, 511 512 ROESY and HRESIMS spectra for compounds 1-11 (PDF). 513 **AUTHOR INFORMATION** 514 515 **Corresponding Authors** *E-mail: lijun9593@gxnu.edu.cn. 516 *E-mail: huangxishan13@foxmail.com. 517 ORCID 518 519 Jun Li: 0000-0003-2023-8609 520 Notes The authors declare no competing financial interest. 521 522

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Captions of Figures



Figure 1. Structures of compounds 1–18 isolated from the leaves of C. paliurus



Figure 2. Key HMBC and COSY correlations of compounds 1–11



Figure 3. ROESY correlations of compounds 1, 2, 4, 7, 10, and 11



Figure 4. Inhibitory effects against NO of the different fractions in LPS-activated RAW 264.7 cells



Figure 5. Inhibitory activities of compounds **7-8** and **10-11** on TNF- α expression in LPS-activated RAW 264.7 cells. The cells were activated with LPS (100 ng/mL) for 2 h and then treated with the compounds **7-8** and **10-11** (10, 20, and 40 μ M) for 24 h. The TNF- α concentration in the supernatants was measured by ELISA kits. Dexamethasone was used as a positive control at the same concentration as the compounds. Values are the mean \pm SD of three independent tests. ### p < 0.001 compared with the control that was induced without LPS. ** p < 0.01 compared with LPS.



Figure 6. Effects of compounds **7-8** and **10-11** on PGE₂ expression by LPS-activated RAW 264.7 cells. The cells were activated with LPS (100 ng/mL) for 2 h and then treated with the compounds **7-8** and **10-11** (10, 20, and 40 μ M) for 24 h. The PGE₂ concentration in the supernatants was determined by ELISA kits. Dexamethasone was used as a positive control at the same concentration as the compounds. Values are the mean \pm SD of three independent tests. ### p < 0.001 compared with the control that was induced without LPS. ** p < 0.01 compared with cells treated with LPS.



Figure 7. Effects of compounds **7-8** and **10-11** on IL-6 expression by LPS-activated RAW 264.7 cells. The cells were activated with LPS (100 ng/mL) for 2 h and then treated with the compounds **7-8** and **10-11** (10, 20, and 40 μ M) for 24 h. The IL-6 concentration in the supernatants was tested by ELISA kits. Dexamethasone was used as a positive control at the same concentration as the compounds. Values are the mean \pm SD of three independent tests. ### p < 0.001 compared with the control that was induced without LPS. ** p < 0.01 compared with cells treated with LPS.



Figure 8. Effects of compound 7 at concentrations of 5, 10, and 20 μ M on iNOS, COX-2, and NF- κ B/p65 expression in LPS-activated RAW 264.7 cells. β -Actin served as the loading control. ### p < 0.001 compared with the control that was activated without LPS. ** p < 0.01 compared with cells treated with LPS.

Captions of Tables

- **Table 1**. ¹H NMR Data for Compounds **1–6** (500 MHz, Pyridine- d_5 , δ in ppm)
- **Table 2**. ¹H NMR Data for Compounds 7–11 (500 MHz, Pyridine- d_5 , δ in ppm)
- **Table 3**. ¹³C NMR Data for Compounds 1–11 (125 MHz, Pyridine- d_5 , δ in ppm)
- Table 4. Inhibitory Activities on NO of Compounds in LPS-activated RAW 264.7

 Cells

Table of Contents Graphic

