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Formosins A–F: Diterpenoids with Anti-microbial Activities from *Excoecaria formosana*



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Abstract Three new halimane-type diterpenoids formosins A–C (1–3), and three clerodane-type diterpenoids formosins D–F (4–6), were isolated from the twigs of *Excoecaria formosana*. Their structures were assigned on the basis of spectroscopic data analysis. Compounds 1 and 4 showed moderate anti-microbial activities against *Bacillus subtilis* (MIC = 50 and 50 µg/mL, respectively). Compound 6 exhibited moderate anti-microbial activities against two strains of *Helicobacter pylori* (*Hp*-SS1 and ATCC 43504) with MIC values of 50 and 50 µg/mL, respectively. *Graphical abstract*



Keywords Excoecaria formosana · Halimane-type · Clerodane-type · Diterpenoid · Anti-microbial

Bing-Dong Lin and Bin Zhou have contributed equally to this work.

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1 Introduction

The genus *Excoecaria* (Euphorbiaceae) compromising 40 species, are widely distributed in Africa and East Asia [1]. Several plants in this genus have been used in folk medicine to treat psoriasis, dermatitis and pruritus [2–4]. The characteristic of this plant genus is the poisonous milk latex, which causes skin blister [5]. Chemical investigations on this plant

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Position	1 (mult., J in Hz)	2 (mult., J in Hz)	3 (mult., J in Hz)	4 (mult., J in Hz)	5 (mult., J in Hz)	6 (mult., <i>J</i> in Hz)
1	α 1.92 (m)	α 1.86 (m)	α 1.90 (m)	α 1.76 (m)	α 1.67 (m)	α 2.89 (dd, 17.9, 14.5)
	β 1.68 (m)	β 1.64 (m)	β 1.68 (m)	β 2.22 (m)	β 2.32 (m)	β 2.66 (dd, 17.9, 3.1)
2	1.64 (m)	1.60 (m)	1.60 (m)	3.61 (m)	4.19 (m)	
3	α 1.36 (td, 12.5,	α 1.35 (td, 12.1,	α 1.36 (m)	5.43 (d, 4.7)	5.27 (br s)	5.78 (s)
	5.4)	4.3)				
	β 1.50 (m)	β 1.46 (m)	β 1.52 (m)			
6	α 2.12 (m)	α 2.12 (m)	α 2.16 (m)	α 1.79 (m)	α 1.80 (m)	α 2.16 (m)
	β 1.94 (m)	β 1.92 (m)	β 1.94 (m)	β 1.29 (m)	β 1.21 (m)	β 1.40 (m)
7	α 1.52 (m)	α 1.52 (m)	α 1.56 (m)	α 2.11 (m)	α 2.11 (m)	α 2.24 (m)
	β 1.70 (m)	β 1.66 (m)	β 1.72 (m)	β 1.47 (m)	β 1.47 (m)	β 1.56 (m)
8	1.78 (m)	1.73 (m)	1.75 (m)	1.61 (m)	1.55 (m)	1.62 (m)
10				1.95 (m)	1.62 (m)	1.90 (m)
11	1.98 (m)	1.98 (m)	2.00 (m)	1.94 (m)	1.92 (m)	1.94 (m)
				2.23 (m)	2.25 (m)	2.18 (m)
12	2.10 (m)	2.02 (m)	2.03 (m)	2.32 (m)	2.34 (m)	2.33 (m)
	2.27 (m)	2.15 (m)	2.18 (m)	2.57 (m)		
14	6.27 (br s)	7.17 (t, 1.6)	6.83 (br s)	6.28 (br s)	6.28 (br s)	6.26 (br s)
15	7.34 (br s)	4.77 (d, 1.8)	5.74 (br s)	7.34 (br s)	7.35 (br s)	7.34 (br s)
16	7.22 (br s)			7.22 (br s)	7.23 (br s)	7.22 (br s)
17	0.95 (d, 6.7)	0.94 (d, 6.7)	0.94 (d, 5.6)	1.16 (d, 6.8)	1.15 (d, 6.9)	1.17 (d, 6.8)
18	1.04 (s)	1.04 (s)	1.03 (s)	1.65 (s)	1.62 (s)	1.90 (s)
19	1.00 (s)	1.00 (s)	0.99 (s)	0.91 (s)	0.99 (s)	1.08 (s)
2-OMe				3.34 (s)		
15-OMe			3.49 (s)			

Table 1 ¹H NMR data for compounds 1–6 in CDCl₃ at 400 MHz

genus have led to the isolation of structurally diverse compounds with significant biological activities including antitumor promoting, anti-ulcer, and anti-microbial activities [6–10]. In the current study, three new halimane-type diterpenoids formosins A–C (1–3), and three clerodane-type diterpenoids formosins D–F (4–6), were isolated from the twigs of *Excoecaria formosana*. Presented herein are the isolation, structural characterization, and biological evaluation of these compounds.

2 Results and Discussion

Compound 1, a white powder, gave a molecular formula $C_{20}H_{28}O_3$ as determined by the (+)-HRESIMS ion at m/z 339.1937 [M + Na]⁺ (calcd 339.1931) requiring seven degrees of unsaturation. The IR absorptions (3000–2800 cm⁻¹, broad band) and (1695 cm⁻¹) showed the presence of a carboxylic group. The diagnostic NMR data (Tables 1, 2) suggested the presence of a β -substituted furan ring (δ_H 6.27, 7.22, and 7.34), a persubstituted double bond, and a carboxylic group (δ_C 181.5). These functionalities

accounted for five out of the seven indices of hydrogen deficiency, requiring the presence of two additional rings in the structure of 1. The aforementioned data suggested that compound 1 is a halimane-type diterpenoid and is structurally related to crotohalimaneic acid [11]. The planar structure of 1 was deduced by 2D NMR spectra. In the HMBC spectrum (Fig. 1), two tertiary methyls ($\delta_{\rm H}$ 1.00, and 1.04, each 3H, s) correlating with C-3, C-4 and C-5 were attached to C-4. The multiple HMBC correlations of H-3, H-7, and H₃-19/C-5 ($\delta_{\rm C}$ 141.3); H-1, H-8, and H-11/C-10 ($\delta_{\rm C}$ 125.2); and H₃-17/C-7, C-8, and C-9 indicated the presence of a typical $\Delta^{5(10)}$ double bond and 8-Me (C-17). The β substituted furan ring and carboxylic group were located at C-12 and C-9 by the HMBC correlations of H-12/C-13, C-14 and C-16; and H-8 and H-11/C-20, respectively. The relative configuration of 1 was established by the ROESY experiment (Fig. 1). The ROESY cross-peaks of H₃-18/H-2 β , H-2 β /H-1 β , H-1 β /H-11, and H-12/H-8 indicated that they are co-facial and randomly assigned in a β -configuration. In consequence, H₃-19 and H₃-17 were α -oriented by the ROESY correlations of H₃-19/H-6 α and H₃-17/H-7 α . The structure of 1 (formosin A) was herein elucidated as shown.

Tabla 2	^{13}C NMP	data fo	r compounde	1 6 in	CDCI	ot 100 MUz	
rame z		α_{a1a}	W COURTAINT	I _ D 10	1 1 1 1 2		

Carbons	1	2	3	4	5	6
1	19.5	19.4	19.4	24.6	30.7	36.5
2	28.2	28.0	28.1	74.3	69.4	199.9
3	39.4	39.3	39.3	120.6	125.1	125.8
4	34.8	34.8	34.8	149.2	146.5	171.4
5	141.3	141.7	142.2	39.3	39.2	40.2
6	25.1	25.0	25.0	36.8	37.1	36.2
7	27.5	27.4	27.4	27.1	27.0	26.7
8	33.6	33.4	33.5	36.8	36.9	36.7
9	54.9	54.6	54.6	49.5	49.4	49.8
10	125.2	125.3	125.1	42.5	46.1	46.1
11	31.0	28.7	28.4 (28.5)	33.7	33.7	33.4
12	18.8	19.5	19.5	17.2	17.4	17.8
13	125.7	134.5	138.9	124.8	124.3	123.9
14	111.0	143.9	141.5	110.9	110.8	110.7
15	142.7	70.2	102.5	142.8	142.9	142.9
16	138.5	174.3	171.3	138.5	138.6	138.7
17	17.4	17.4	17.4	16.5	16.6	16.3
18	26.6	26.6	26.6	18.1	17.9	19.2
19	29.0	28.9	28.9	16.0	17.4	16.0
20	181.5	180.4	180.0	182.6	181.8	181.3
2-OMe				56.3		
15-OMe			56.9 (57.0)			



Fig. 1 a Selected HMBC, and b ROESY correlations of 1



Fig. 2 a Selected HMBC, and b ROESY correlations of 4

Compound **2** had the molecular formula of $C_{20}H_{28}O_4$, as determined by the ¹³C NMR data and the (+)-HRESIMS ion at m/z 355.1839 [M + Na]⁺ (calcd 355.1880), which is 16 mass units more than that of **1**. Comparison of its NMR spectroscopic data (Tables 1, 2) with those of **1** revealed they are structural analogues with the obvious difference being the presence of an α,β -unsaturated- γ -lactone moiety instead of the β -substituted furan ring. It was confirmed by the NMR data (δ_H 4.77 and 7.17; δ_C 70.2, 134.5, 143.9, and 174.3), as well as the key HMBC correlations from H-12 to C-13, C-14, and C-16 (Figure S13, Supporting Information). Thus, the structure of **2** (formosin B) was determined as shown.

Compound **3** displayed a molecular formula of $C_{21}H_{30}O_5$ as established by the (+)-HRESIMS at m/z 385.1991 [M + Na]⁺ (calcd 385.1991) and the ¹³C NMR data. Analysis of the NMR data (Tables 1, 2) of **3** showed many similarities to those of **2**. The only difference was the presence of an additional methoxy group (δ_H 3.49, s, 3H), which was located at C-15 to form the acetal motif, which was confirmed by the downfield shifted C-15 ($\Delta \delta_C$ 32.3) as compared to that of **2**. Compound **3** was obtained as a pair of inseparable C-15 epimers, which exhibited several pairs of very close carbon resonances in the ¹³C NMR spectrum (Figure S19, Supplementary Material). Therefore, the

structure of **3** (formosin C) with its relative configuration was confirmed as depicted by the HMBC and ROESY spectra (Figures S21 and S22, Supplementary Material).

Compound 4 was obtained as a white powder with a molecular formula of $C_{21}H_{30}O_4$ as established by the ¹³C NMR data and the (+)-HRESIMS ion at m/z 369.1970 $[M + Na]^+$ (calcd 369.2036), demanding seven degrees of unsaturation. The IR absorption bands $(3000-2800 \text{ cm}^{-1})$, broad band; and 1695 cm⁻¹) showed the presence of a carboxylic group. The characteristic NMR signals for a β furan ring, a trisubstituted double bond, a methoxyl and a carboxylic groups were observed from the ¹H and ¹³C NMR spectroscopic data analysis (Tables 1, 2). Comprehensive analysis of the NMR spectra of 4 revealed its structure is highly related with that of junceic acid [12] with a clerodane-type diterpenoid backbone. The only difference was the presence of an additional methoxyl group in 4, which was placed at C-2 by the HMBC correlation (Fig. 2) from CH₃O ($\delta_{\rm H}$ 3.34) to C-2 ($\delta_{\rm C}$ 74.3). The carboxylic group was attached to C-9 via the HMBC correlations from H-8, H-10 and H-11 to C-20 ($\delta_{\rm C}$ 182.6). The key HMBC cross-peaks from H₃-18 to C-3 ($\delta_{\rm C}$ 120.6), C-4 ($\delta_{\rm C}$ 149.2) and C-5, and from H₃-19 to C-4 revealed the presence of Δ^3 double bond. The relative configuration of 4 was established by the ROESY experiment (Fig. 2). The ROESY correlations of H-10/H-6 β and H-10/H-8 indicated that H-8 and H-10 are co-facial and randomly

assigned in a β -configuration. Consequently, H₃-19 and H-2 were thus assigned to be α -directed by the ROESY correlations of H₃-19/H-1 α , H₃-19/H-7 α , and H-2/H-1 α . Therefore, the structure of **4** (formosin D) was established as depicted.

Compound **5** possessed a molecular formula $C_{20}H_{28}O_4$ based on the ¹³C NMR data and the (+)-HRESIMS ion at *m*/z 355.1885 [M + Na]⁺ (calcd 355.1888), which is 14 mass units less than that of **4**. Detailed analysis of the NMR data (Tables 1, 2) of **5** revealed that its structure is closely related with that of **4** with the only difference being the absence of the methyl etherification, which is consistent with the molecular formula. The structure of **5** (formosin E) with the relative configuration was further confirmed by HMBC and ROESY spectra (Figures S39 and S40, Supplementary Material).

Compound **6**, named formosin F, exhibited a sodiated molecular ion at m/z 353.1724 $[M + Na]^+$ (calcd 353.1723) in the (+)-HRESIMS, consistent with a molecular formula of $C_{20}H_{26}O_4$, which was supported by the ¹³C NMR data. Comparison of the NMR data (Tables 1, 2) of **6** with those of **5** revealed they are structural analogues. The main difference was the presence of a keto group at C-2 in **6** instead of the hydroxy group in the latter, which was confirmed by the HMBC correlations (Figure S42, Supporting Information) from H-3 and H-10 to C-2 (δ_C 199.9). The structure of **6** was thus determined as shown.

The new isolates were tested for anti-microbial activities against a panel of microbes *in vitro* by the microdilution method [13, 14]. Compounds 1 and 4 exhibited moderate activity against *Bacillus subtilis ATCC 6633* with MIC values of 50 and 50 µg/mL, respectively, where magnolol was used as the positive control (MIC = 12.5μ g/mL). Compound 6 showed moderate antibacterial activities against two strains *Helicobacter pylori* (*Hp*-SS1 or ATCC 43504) with MIC values of 50 and 50 µg/mL, respectively, and metronidazole was used as the positive control (MIC = 0.312 and 128μ g/mL, respectively).

3 Experimental Section

3.1 General Experimental Procedures

Optical rotations were determined on a Perkin-Elmer 341 polarimeter. UV spectra were recorded on a Shimadzu UV-2550 spectrophotometer. IR spectra were acquired on a Perkin-Elmer 577 spectrometer. NMR spectra were measured on a Bruker AM-400 spectrometer with TMS as internal standard. ESIMS and HRESIMS were performed on a Bruker Daltonics Esquire 3000 plus and a Waters-Micromass Q-TQF Ultima Global mass spectrometer, respectively. Semi-preparative HPLC was performed on a

Waters 1525 binary pump system with a Waters 2489 detector (210 nm) and equipped with a YMC-Pack ODS-A (250 \times 10 mm, S-5 μ m). Silica gel (200–300 mesh, Qingdao Haiyang Chemical Co., Ltd.), C₁₈ reversed-phase (RP-18) silica gel (20–45 μ m, Fuji Silysia Chemical Ltd.), CHP20P MCI gel (75–150 μ m, Mitsubishi Chemical Corporation), and Sephadex LH-20 gel (Amersham Biosciences) were used for column chromatography (CC). Precoated silica gel GF₂₅₄ plates (Qingdao Haiyang Chemical Co., Ltd.) were used for TLC detection. All solvents used for CC were of analytical grade (Shanghai Chemical Reagents Co., Ltd.), and solvents used for HPLC were of HPLC grade (J & K Scientific Ltd.).

3.2 Plant Material

The twigs of *E. formosana* were collected from Sanya city of Hainan Province, the People's Republic of China, and authenticated by Prof. S.-M. Huang, Department of Biology, Hainan University. A voucher specimen has been deposited in Shanghai Institute of Materia Medica, Chinese Academy of Sciences (accession number: SMEF-2006-1Y).

3.3 Extraction and Isolation

The air-dried, powdered twigs of E. formosana (6.0 kg) was extracted three times with 95 % EtOH at room temperature to give a crude extract (290 g), which was partitioned between EtOAc and H2O. The EtOAc-soluble fraction (85 g) was subjected to passage over an MCI gel column (MeOH/H₂O, 3:7-9:1) to afford fractions A-G. Fraction C (25.7 g) was separated over a silica gel column eluted with gradient mixtures of petroleum ether-acetone (35:1-1:1, v/v) to afford major fractions C1-C6. Fraction C3 (3.4 g) was separated on a reversed-phase C_{18} silica gel column (MeOH/H₂O, 55-100 %) to yield three major portions (C3a-C3c), and each of those was purified by a semi-preparative HPLC (60 % CH₃CN in H₂O, 3 mL/min) to yield compounds 1 (20 mg), 2 (10 mg), and 6 (100 mg), respectively. Fraction C4 (515 mg) was purified by a semipreparative HPLC (55 % CH₃CN in H₂O, 3 mL/min) to give compound 4 (15 mg). Fraction C6 (1.5 g) was separated on a column of Sephadex LH-20, and then purified by a semi-preparative HPLC (50 % CH₃CN in H₂O, 3 mL/ min) to yield compound 5 (8 mg). Fraction E (11.4 g) was chromatographed on a silica gel column eluted with petroleum ether-ethyl acetate (25:1-1:4, v/v) to afford subfractions E1-E4. Fraction E2 (217 mg) was separated on a reversed-phase column containing C₁₈ silica gel (MeOH/ H_2O , 70–100%) to yield three fractions E2a–E2c. Fraction E2b (35 mg) was separated by a semi-preparative HPLC (70 % CH₃CN in H₂O, 3 mL/min) to yield compound **3** (8 mg).

3.4 Formosin A (1)

White powder; $[\alpha]_D^{21}$ +123 (*c* 0.10, MeOH); UV (MeOH) λ_{max} (log ε) 206 (4.19) nm; IR (KBr) v_{max} 3444, 3000–2500, 1695, 1622, 1458, 1257, 1024, 600 cm⁻¹; ¹H NMR (CDCl₃), see Table 1 and ¹³C NMR (CDCl₃) see Table 2; (+)-ESIMS *m*/*z* 339.2 [M + Na]⁺; (+)-HRE-SIMS *m*/*z* 339.1937 [M + Na]⁺ (calcd for C₂₀H₂₈O₃Na, 339.1931).

3.5 Formosin B (2)

White powder; $[\alpha]_D^{21}$ +212 (*c* 0.10, MeOH); UV (MeOH) λ_{max} (log ε) 207 (3.92); IR (KBr) v_{max} 3435, 3000–2500, 1753, 1695, 1456, 1205, 1070, 756 cm⁻¹; ¹H NMR (CDCl₃), see Table 1 and ¹³C NMR (CDCl₃) see Table 2; (+)-ESIMS *m*/*z* 355.2 [M + Na]⁺; (+)-HRESIMS *m*/*z* 355.1839 [M + Na]⁺ (calcd for C₂₀H₂₈O₄Na, 355.1880).

3.6 Formosin C (3)

White powder; $[\alpha]_D^{21} + 22$ (*c* 0.10, MeOH); UV (MeOH) λ_{max} (log ε) 211 (3.58); IR (KBr) v_{max} 3425, 3000–2500, 1726, 1693, 1659, 1468, 1178, 1065, 600 cm⁻¹; ¹H NMR (CDCl₃), see Table 1 and ¹³C NMR (CDCl₃) see Table 2; (+)-ESIMS *m*/*z* 385.2 [M + Na]⁺; (+)-HRESIMS *m*/*z* 385.1991 [M + Na]⁺ (calcd for C₂₁H₃₀O₅Na, 385.1991).

3.7 Formosin D (4)

White powder; $[\alpha]_D^{21} - 164$ (*c* 0.025, MeOH); UV (MeOH) λ_{max} (log ε) 204 (4.21); IR (KBr) v_{max} 3427, 3000–2500, 1695, 1452, 1254, 1082 cm⁻¹; ¹H NMR (CDCl₃), see Table 1 and ¹³C NMR (CDCl₃) see Table 2; (+)-ESIMS *m*/*z* 369.2 [M + Na]⁺; (+)-HRESIMS *m*/*z* 369.1970 [M + Na]⁺ (calcd for C₂₁H₃₀O₄Na, 369.2036).

3.8 Formosin E (5)

White powder; $[\alpha]_D^{21} -20$ (*c* 0.05, MeOH); UV (MeOH) λ_{max} (log ε) 199 (3.89); IR (KBr) ν_{max} 3419, 3000–2500, 1695, 1452, 1383, 1244, 1026, 874, 600 cm⁻¹; ¹H NMR (CDCl₃), see Table 1 and ¹³C NMR (CDCl₃) see Table 2; (+)-ESIMS *m*/*z* 355.2 [M + Na]⁺; (+)-HRESIMS *m*/*z* 355.1885 [M + Na]⁺ (calcd for C₂₀H₂₈O₄Na, 355.1888).

3.9 Formosin F (6)

White powder; $[\alpha]_D^{21} - 175$ (*c* 0.10, MeOH); UV (MeOH) λ_{max} (log ε) 237 (4.86); IR (KBr) v_{max} 3000–2500, 1703, 1633, 1435, 1381, 1219, 1026, 874 cm⁻¹; ¹H NMR (CDCl₃), see Table 1 and ¹³C NMR (CDCl₃) see Table 2; (+)-ESIMS m/z 331 [M + H]⁺; (+)-HRESIMS m/z 353.1724 [M + Na]⁺ (calcd for C₂₀H₂₆O₄Na, 353.1723).

3.10 Antimicrobial Activity Assay

The in vitro antibacterial activities against *Bacillus subtilis* ATCC 6633 were tested by applying the protocols described in our previous research [13].

Antibacterial tests against *Helicobacter pylori* strains (*Hp*-SS1 or ATCC 43504 strain) were carried out in vitro according to the protocols described previously [14].

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Compliance with Ethical Standards

Conflict of Interest The authors declare no conflict of interest.

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References

- J.S. Ma, Zhongguo Zhiwu Zhi, vol. 44 (Science Press, Beijing, 1997), p. 8
- P.M. Giang, P.T. Son, K. Matsunami, H. Otsuka, Chem. Pharm. Bull. 53, 1600–1603 (2005)
- D.S. Ning, L.Y. Peng, S.H. Lv, D.P. Li, Z.H. Pan, Nat. Prod. Res. 29, 524–528 (2015)
- Z.C. Wang, Y.M. Lin, D.Q. Feng, C.H. Ke, P. Lin, C.L. Yan, J.D. Chen, Molecules 14, 414–422 (2009)
- S. Kumarasinghe, R. Seneviratne, Australas. J. Dermatol. 39, 275–276 (1998)
- T. Konishi, K. Yamazoe, T. Konoshima, T. Maoka, Y. Fujiwara, K. Miyahara, J. Nat. Prod. 66, 108–111 (2003)
- T. Konishi, T. Konoshima, Y. Fujiwara, S. Kiyosawa, J. Nat. Prod. 63, 344–346 (2000)
- G. Agoramoorthy, M. Chandrasekaran, V. Venkatesalu, M. Hsu, Braz. J. Microbiol. 38, 739–742 (2007)
- P. Thirunavukkarasu, L. Ramkumar, T. Ramanathan, Global. J. Pharmacol. 3, 123–126 (2009)
- T. Konishi, M. Takasaki, H. Tokuda, S. Kiyosawa, T. Konoshima, Biol. Pharm. Bull. 21, 993–996 (1998)
- S. Roengsumran, S. Pornpakakul, N. Muangsin, P. Sangvanich, T. Nhujak, P. Singtothong, N. Chaichit, S. Puthong, A. Petsom, Planta Med. **70**, 87–89 (2004)
- Y. Asakawa, M. Toyota, A. Ueda, Phytochemistry 29, 2165–2167 (1990)
- S.P. Yang, J.M. Yue, Bioorg. Med. Chem. Lett. 11, 3119–3122 (2001)
- S. Yin, C.Q. Fan, L. Dong, J.M. Yue, Tetrahedron 62, 2569–2575 (2006)