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## Molecular and Crystal Structures of $\alpha,\alpha,\beta$ -Trimethylfuranylxanthone from *Cratoxylum formosum* ssp. *pruniflorum*: A Partial Racemate

NAWONG BOONNAK,<sup>1,\*</sup> SUCHADA CHANTRAPROMMA,<sup>2,\*</sup>  
AND HOONG-KUN FUN<sup>3,4</sup>

<sup>1</sup>Department of Chemistry, Faculty of Science, Thaksin University, Pattalung, Thailand

<sup>2</sup>Faculty of Science, Department of Chemistry, Prince of Songkla University, Hat-Yai, Songkhla, Thailand

<sup>3</sup>X-ray Crystallography Unit, School of Physics, Universiti Sains Malaysia, USM, Penang, Malaysia

<sup>4</sup>Department of Pharmaceutical Chemistry, College of Pharmacy, King Saud University, Riyadh, Saudi Arabia

*The title xanthone known as Formoxanthone C (1) was isolated from the green fruits of Cratoxylum formosum ssp. pruniflorum. Compound 1 crystallized out in P2<sub>1</sub>/c space group, indicating that 1 exists both enantiomers. The unit cell parameters are a = 13.8038(2), b = 14.6712(2), c = 9.4686(1) Å,  $\beta$  = 95.358(1)°, V = 1909.19(4) Å<sup>3</sup>, and Z = 4. The  $\alpha,\alpha,\beta$ -trimethyldihydrofuran ring was in an envelope conformation and disordered over two sites with 0.628(1) and 0.372(1) occupancies for the major and minor components, respectively, which corresponded to the less furan ring strain and the steric hindrance between the two methyl substituents (at positions 1' and 2') on the furan ring of the major component than that of the minor component. The crystal structure is stabilized by intermolecular O–H...O hydrogen bonds and weak C–H...O interactions.*

**Keywords** Crystal structure; formoxanthone C;  $\alpha,\alpha,\beta$ -trimethyldihydrofuran; partial racemate; xanthone; X-ray

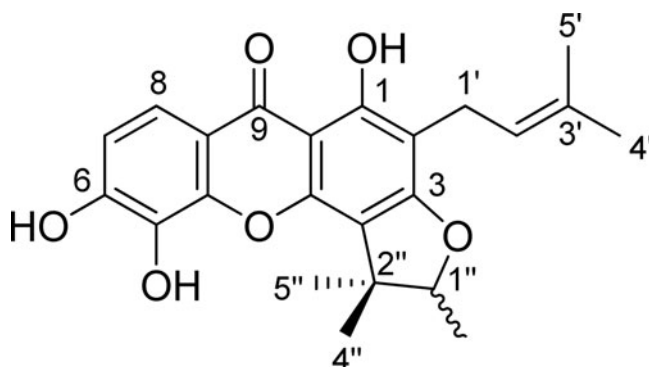
### Introduction

*Cratoxylum formosum* ssp. *pruniflorum*, belonging to the family Guttiferae, is widely distributed in northern and north-eastern parts of Thailand [1]. Decoction of the fresh or dried leaves from this plant has been used in folk medicine as stomach and diuretic tonics [2]. Previous reports [3–5] have revealed that tri- and tetra-oxygenated xanthones were mainly produced as major components from plants of *Cratoxylum* genus. Moreover, several

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\*Address correspondence to Nawong Boonnak and Suchada Chantrapromma, Faculty of Traditional Thai Medicine, Prince of Songkla University, Hat-Yai, Songkhla 90112, Thailand and Faculty of Science, Department of Chemistry, Prince of Songkla University, Hat-Yai, Songkhla 90112, Thailand. E-mail: suchada.c@psu.ac.th; nawongb@yahoo.com

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**Figure 1.** Schematic diagram of formoxanthone C (**1**).

isolated xanthenes were reported to possess various interesting biological activities [4, 6–8]. The bioactivity-guided fractionation of the  $\text{CH}_2\text{Cl}_2$  crude extract of *C. formosum* spp. *pruniflorum* green fruits afforded compound **1** (Fig. 1), which known as formoxanthone C [6] [systematic name: 5,9,10-trihydroxy-1,1,2-trimethyl-4-(3-methylbut-2-enyl)-1*H*-furo[2,3-*c*]xanthene-6(2*H*)-one]. Our nitric oxide inhibitory activity study resulted that compound **1** exhibits potent nitric oxide inhibitory activity with  $\text{IC}_{50}$  value of  $4.3 \mu\text{M}$  [9] and also showed strong antibacterial activity against methicillin-resistant *Staphylococcus aureus* (MRSA) ATCC43300, *Enterococcus faecalis* TISTR459, vancomycin-resistant *Enterococcus faecalis* (VRE) ATCC51299, *Bacillus subtilis*, and *Salmonella typhi* with an MIC value of  $4.67 \mu\text{g/mL}^{-1}$  [9]. It is well documented that bioactivities are related to the structure of the compound thereby making the correct characterization and identification of compounds of paramount importance. Of all the various physical methods that are important in helping to elucidate structure, single crystal X-ray diffraction is a powerful technique and undoubtedly the most far reaching and complete method. To study in more details of the molecular structure of **1** and to establish the conformation of the  $\alpha,\alpha,\beta$ -trimethyldihydrofuran ring, its molecular and crystal structures by single crystal X-ray diffraction was studied. The results reported here show that **1** exists in both enantiomers and also in two conformations with 62.8% for the major and 37.2% for the minor components.

## Experimental

### General Experimental Procedures

Melting points were determined with a Fisher–John melting point apparatus. Optical rotations were measured on a JASCO P-1020 digital polarimeter. UV-Vis and IR spectra were recorded on SPECORD S 100 (Analytikjena) and Perkin–Elmer FTS FT-IR spectrophotometer, respectively. The  $^1\text{H}$  and  $^{13}\text{C}$  NMR spectra were recorded on a 300 MHz Bruker FT-NMR Ultra Shield™ spectrometer in  $\text{CDCl}_3$  with TMS as an internal standard. Chemical shifts are reported in  $\delta$  (ppm) and coupling constants ( $J$ ) are expressed in hertz. Quick column chromatography (QCC) and column chromatography (CC) were carried out on silica gel 60 F<sub>254</sub> (Merck) and silica gel 100 (Merck), respectively.

### Extraction and Isolation

Air-dried green fruits of *C. formosum* ssp. *pruniflorum* (5.0 kg) were extracted with  $\text{CH}_2\text{Cl}_2$  and  $\text{CH}_3\text{OH}$  (each  $2 \times 20 \text{ L}$ , for a week) successively at room temperature and further

evaporated under reduced pressure to afford the crude extracts of  $\text{CH}_2\text{Cl}_2$  (31.42 g) and  $\text{CH}_3\text{OH}$  (65.74 g), respectively. The crude  $\text{CH}_2\text{Cl}_2$  extract (31.42 g) of the green fruits of *C. formosum* ssp. *pruniflorum* was further subjected to QCC on silica gel using hexane as a first eluent and then increasing the polarity with acetone to give 14 fractions (F1–F14). Fraction F10 was separated by QCC eluting with a gradient of acetone-hexane to give 17 subfractions (F10A–F10Q). Subfraction F10N was separated by CC and eluted with a gradient of EtOAc-hexane to give eight subfractions (F10N1–F10N8). Subfraction F10N6 was separated by CC and eluted with  $\text{CHCl}_3$  to give the compound **1**. Yield: yellow solid 15.2 mg, m.p.: 152–154°C;  $[\alpha]_D^{25} = -44.8$  ( $c$  0.05,  $\text{CHCl}_3$ ); UV-Vis ( $\text{CH}_3\text{OH}$ )  $\lambda_{\text{max}}$  (log  $\epsilon$ ) 258 (4.51), 276 (4.44), 392 (3.85) nm; FT-IR (KBr)  $\nu_{\text{max}}$  3440, 1646, 1624  $\text{cm}^{-1}$ ;  $^1\text{H}$  NMR ( $\text{CD}_3\text{OD}+\text{CDCl}_3$ , 300 MHz, ppm)  $\delta$ : 13.40 (1H, s, OH), 7.63 (1H, d,  $J = 8.1$  Hz, H-8), 6.85 (1H, br s, H-7), 5.28 (1H, br t,  $J = 7.2$  Hz, H-2'), 4.52 (1H, q,  $J = 6.6$  Hz, H-1''), 3.29 (2H, d,  $J = 7.2$  Hz,  $\text{CH}_2$ -1'), 1.77 (3H, s,  $\text{CH}_3$ -5'), 1.68 (3H, s,  $\text{CH}_3$ -4'), 1.59 (3H, s,  $\text{CH}_3$ -5''), 1.41 (3H, dd,  $J = 6.6, 1.2$  Hz,  $\text{CH}_3$ -3''), and 1.31 (3H, s,  $\text{CH}_3$ -4'').  $^{13}\text{C}$  NMR ( $\text{CD}_3\text{OD}+\text{CDCl}_3$ , 75 MHz)  $\delta$ : 164.5 (C-1), 110.6 (C-2), 168.0 (C-3), 115.9 (C-4), 135.9 (C-5), 149.7 (C-6), 115.9 (C-7), 121.1 (C-8), 184.5 (C-9), 154.9 (C-4a), 154.1 (C-5a), 118.2 (C-8a), 106.8 (C-9a), 25.6 (C-1'), 125.6 (C-2'), 135.9 (C-3'), 29.6 (C-4'), 21.6 (C-5'), 44.4 (C-1''), 47.9 (C-2''), 18.2 (C-3''), 25.1 (C-4''), and 29.6 (C-5'').

### Preparation of Single Crystal for Structure Determination

Yellow block-shaped single crystals of formoxanthone **C** suitable for X-ray diffraction study were obtained from  $\text{CH}_2\text{Cl}_2$ – $\text{CH}_3\text{OH}$  (4:1 v/v) solution by slow evaporation at room temperature after several days. A suitable single crystal was selected optically for the diffraction study.

A single crystal of dimensions 0.35 mm  $\times$  0.30 mm  $\times$  0.17 mm was used. Crystallographic data were collected at 100.0 (1) K with the Oxford Cryosystem Cobra low-temperature attachment. The data were collected using a Bruker Apex2 CCD diffractometer with a graphite monochromated  $\text{Mo K}\alpha$  ( $\lambda = 0.71073$  Å) radiation at a detector distance of 5 cm using APEX2 [10]. The collected data were reduced using SAINT program [10], and the empirical absorption corrections were performed using SADABS program [10]. The structures were solved by direct methods and refined by least-squares using the SHELXTL software package [11]. All nonhydrogen atoms were refined anisotropically. Hydroxy H atoms were located in different maps and refined isotropically, whereas the remaining H atoms were placed in calculated positions with C–H = 0.93 Å for aromatic, 0.98 Å for CH, 0.97 Å for  $\text{CH}_2$ , and 0.96 Å for  $\text{CH}_3$ . The  $U_{\text{iso}}(\text{H})$  values were constrained to be  $1.5U_{\text{eq}}$  of the carrier atoms for methyl H atoms and  $1.2U_{\text{eq}}$  for hydroxyl and the other H atoms. The final refinement converged well. Materials for publication were prepared using SHELXTL [11], PLATON [12], and Mercury [13]. The  $\alpha,\alpha,\beta$ -trimethylfuran ring was found to be disordered over two sites with a 0.628(1):0.372(1) refined site occupancy ratio. The crystal data of formoxanthone **C** (CCDC 910092) are summarized in Table 1.

## Results and Discussion

### Spectroscopic Properties

The UV-Vis spectrum showed absorption bands of a xanthone at 258, 276, and 392 nm. The FT-IR spectrum exhibited the hydroxyl group at 3440  $\text{cm}^{-1}$  and conjugated carbonyl group at 1646  $\text{cm}^{-1}$ . The  $^1\text{H}$  NMR spectrum ( $\text{CD}_3\text{OD}+\text{CDCl}_3$ ) exhibited a chelated hydroxyl proton at  $\delta$  13.40 (1H, s), a pair of *ortho*-coupled aromatic protons at  $\delta$  7.63

**Table 1.** Crystal data and structure refinement for formoxanthone C

CCDC reference no.	CCDC 910092
Empirical formula	C <sub>23</sub> H <sub>24</sub> O <sub>6</sub>
Formula weight	396.43
Crystal system, space group	Monoclinic, <i>P</i> 2 <sub>1</sub> / <i>c</i>
Temperature (K)	100.0(1)
<i>a</i> , <i>b</i> , <i>c</i> (Å)	13.8038(2), 14.6712(2), 9.4686(1)
$\beta$ (°)	95.358(1)
<i>V</i> (Å <sup>3</sup> )	1909.19(4)
<i>Z</i>	4
Calculated density (Mg/m <sup>−3</sup> )	1.379
Radiation type, wavelength (Å)	Mo K $\alpha$ , 0.71073
Absorption coefficient (mm <sup>−1</sup> )	0.098
Crystal form, color	block, yellow
Crystal size (mm)	0.35 × 0.30 × 0.17
Diffractometer	CCD area detector diffractometer
Final <i>R</i> indices [ <i>I</i> > 2 $\sigma$ ( <i>I</i> )]	<i>R</i> <sub>1</sub> = 0.0619, <i>wR</i> <sub>2</sub> = 0.1550
<i>R</i> indices (all data)	<i>R</i> <sub>1</sub> = 0.0804, <i>wR</i> <sub>2</sub> = 0.1655
Reflections collected / unique	29475/4383 [ <i>R</i> (int) = 0.0391]
$\theta$ range for data collection (°)	2.03–27.50
Limiting indices	−17 < = <i>h</i> < = 17, −16 < = <i>k</i> < = 19, −12 < = 1 < = 11
Refinement method	Full-matrix least-squares on <i>F</i> <sup>2</sup>
Data/restraints/parameters	4383/0/298
Completeness to $\theta$ = 27.50	100.0%
<i>F</i> (000)	840
Extinction method	None
Goodness-of-fit on <i>F</i> <sup>2</sup>	1.101
Largest difference in peak and hole (e Å <sup>−3</sup> )	0.782 and −0.325

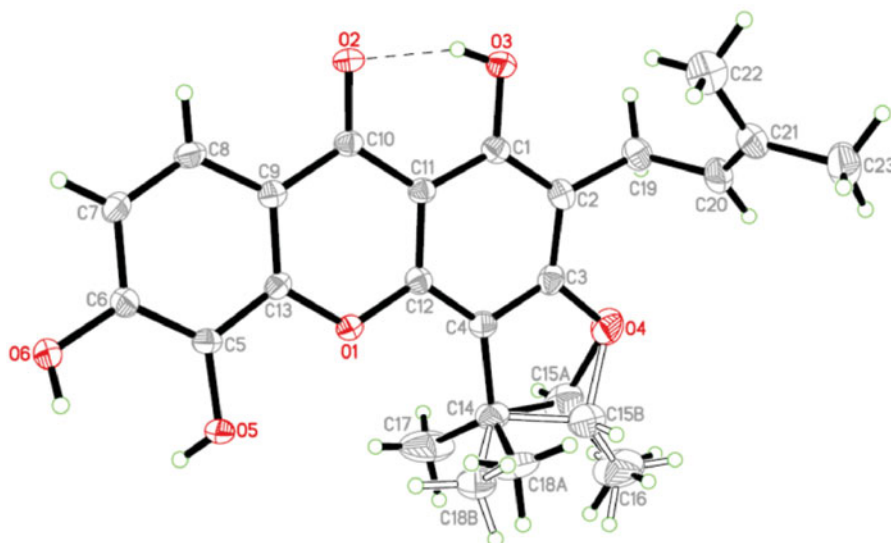
(1H, d, *J* = 8.1 Hz, H-8) and 6.85 (1H, br s, H-7), an isoprenyl side chain at  $\delta$  5.28 (1H, br t, *J* = 7.2 Hz, H-2'), 3.29 (2H, d, *J* = 7.2 Hz, H<sub>2</sub>-1'), 1.77 (3H, s, CH<sub>3</sub>-5'), and 1.68 (3H, s, CH<sub>3</sub>-4'). Moreover, the presence of a  $\alpha,\alpha,\beta$ -trimethyldihydrofuran ring was suggested by the following <sup>1</sup>H NMR spectral data at  $\delta$  4.52 (1H, q, *J* = 6.6 Hz, H-1''), 1.59 (3H, s, CH<sub>3</sub>-5''), 1.41 (3H, dd, *J* = 6.6, 1.2 Hz, CH<sub>3</sub>-3''), and 1.31 (3H, s, CH<sub>3</sub>-4''), respectively.

### Crystal Structure

The X-ray diffraction of compound **1** was studied and revealed its molecular structure, which was shown in Figs. 2 and 3. The chemical structure of **1** showed that it contains one chiral center at C-1'' (Fig. 1). However, compound **1** crystallized out in the centrosymmetric space group *P*2<sub>1</sub>/*c* which clearly indicated that the compound contained both enantiomers [(1'*S*)-**1** and (1'*R*)-**1**] which can be clearly seen in the crystal packing in Fig. 4. Therefore, compound **1** was a racemic mixture. Bond lengths, angles, and torsion angles of **1** are listed in Table 2 and hydrogen bonds are listed in Table 3.

**Table 2.** Selected bond lengths (Å), angles, and torsion angles (°) for **1**

Bond lengths					
O1—C13	1.361(2)	C7—H <sub>7A</sub>	0.93	C16—H <sub>16D</sub>	0.96
O1—C12	1.372(2)	C8—C9	1.406(3)	C16—H <sub>16E</sub>	0.96
O2—C10	1.270(2)	C8—H <sub>8A</sub>	0.9300	C16—H <sub>16F</sub>	0.96
O3—C1	1.350(3)	C9—C13	1.398(3)	C17—H <sub>17A</sub>	0.96
O3—H1 O <sub>3</sub>	0.78(3)	C9—C10	1.459(3)	C17—H <sub>17B</sub>	0.96
O4—C3	1.355(3)	C10—C11	1.435(3)	C17—H <sub>17C</sub>	0.96
O4—C15A	1.499(5)	C11—C12	1.412(3)	C18A—H <sub>18A</sub>	0.96
O4—C15B	1.629(8)	C14—C18A	1.449(6)	C18A—H <sub>18B</sub>	0.96
O5—C5	1.369(2)	C14—C17	1.493(4)	C18A—H <sub>18C</sub>	0.96
O5—H1 O <sub>5</sub>	0.82(3)	C14—C15B	1.566(8)	C18B—H <sub>18D</sub>	0.96
O6—C6	1.361(3)	C8—H <sub>8A</sub>	0.93	C18B—H <sub>18E</sub>	0.96
O6—H1 O <sub>6</sub>	0.84(3)	C9—C13	1.398(3)	C18B—H <sub>18F</sub>	0.96
C1—C2	1.390(3)	C14—C15A	1.628(5)	C19—C20	1.517(3)
C1—C11	1.430(3)	C14—C18B	1.661(10)	C19—H <sub>19A</sub>	0.97
C2—C3	1.394(3)	C15A—C16	1.472(6)	C19—H <sub>19B</sub>	0.97
C2—C19	1.514(3)	C15A—H <sub>15A</sub>	0.98	C20—C21	1.323(4)
C3—C4	1.394(3)	C15A—H <sub>16E</sub>	1.4613	C20—H <sub>20A</sub>	0.93
C4—C12	1.376(3)	C15B—C16	1.426(9)	C21—C23	1.503(4)
C4—C14	1.515(3)	C15B—H <sub>15D</sub>	0.98	C21—C22	1.504(4)
C5—C6	1.391(3)	C15B—H <sub>16B</sub>	1.4382	C22—H <sub>22A</sub>	0.96
C5—C13	1.399(3)	C16—H <sub>16A</sub>	0.96	C22—H <sub>22B</sub>	0.96
C6—C7	1.405(3)	C16—H <sub>16B</sub>	0.96	C22—H <sub>22C</sub>	0.96
C7—C8	1.372(3)	C16—H <sub>16C</sub>	0.96	C23—H <sub>23A</sub>	0.96
Bond angles					
C13—O1—C12	119.47(16)	O6—C6—C7	117.99(19)	C18A—C14—C17	117.9(4)
C3—O4—C15A	105.0(2)	C5—C6—C7	120.26(19)	C18A—C14—C4	110.3(3)
C3—O4—C15B	105.0(3)	C8—C7—C6	120.4(2)	C17—C14—C4	113.6(2)
O3—C1—C2	118.17(19)	C7—C8—C9	120.69(19)	C17—C14—C15B	128.3(5)
O3—C1—C11	119.86(19)	C13—C9—C8	118.15(19)	C18A—C14—C15A	114.9(3)
C2—C1—C11	121.97(19)	C13—C9—C10	119.02(18)	C17—C14—C18B	96.5(4)
C1—C2—C3	115.65(19)	C8—C9—C10	122.83(18)	C4—C14—C18B	113.6(4)
C1—C2—C19	123.0(2)	O2—C10—C11	121.38(19)	C15B—C14—C18B	103.8(5)
C3—C2—C19	121.3(2)	O2—C10—C9	121.81(19)	C15A—C14—C18B	135.0(4)
O4—C3—C2	121.6(2)	C11—C10—C9	116.81(18)	C16—C15A—O4	111.2(4)
O4—C3—C4	112.87(19)	C12—C11—C1	118.06(19)	C16—C15A—C14	116.2(4)
C2—C3—C4	125.6(2)	C12—C11—C10	119.82(19)	O4—C15A—C14	105.2(3)
C12—C4—C3	116.99(19)	C1—C11—C10	122.09(18)	C16—C15B—C14	123.2(7)
C12—C4—C14	131.3(2)	O1—C12—C4	116.63(18)	C16—C15B—O4	106.6(6)
C3—C4—C14	111.64(19)	O1—C12—C11	121.73(18)	C14—C15B—O4	102.1(5)
O5—C5—C6	122.87(18)	C4—C12—C11	121.64(19)	C2—C19—C20	114.59(19)
O5—C5—C13	118.59(18)	O1—C13—C9	122.95(18)	C21—C20—C19	126.8(2)
C6—C5—C13	118.50(18)	O1—C13—C5	115.17(17)	C20—C21—C23	122.2(2)
O6—C6—C5	121.72(19)	C9—C13—C5	121.88(19)	C20—C21—C22	123.6(2)
C13—O1—C12	119.47(16)	O6—C6—C7	117.99(19)	C23—C21—C22	114.2(2)
Torsion angles					
O3—C1—C2—C3	−177.11(18)	C8—C9—C10—O2	2.5(3)	C12—C4—C14—C18B	−51.0(5)
C11—C1—C2—C19	−178.10(19)	O3—C1—C11—C10	0.9(3)	C3—C4—C14—C18B	127.3(5)
C15A—O4—C3—C4	19.2(3)	O2—C10—C11—C1	−0.8(3)	C3—O4—C15A—C16	−156.5(3)
C15B—O4—C3—C4	−17.3(5)	C13—O1—C12—C11	3.0(3)	C18A—C14—C15A—C16	35.8(5)
C1—C2—C3—O4	177.97(19)	C14—C4—C12—O1	1.7(3)	C17—C14—C15A—C16	−92.9(4)
C19—C2—C3—C4	179.0(2)	O5—C5—C13—C9	−175.31(18)	C4—C14—C15A—C16	151.4(4)
O4—C3—C4—C14	0.6(3)	C3—C4—C14—C18A	101.3(4)	C18B—C14—C15B—C16	98.3(9)
C2—C3—C4—C14	−179.7(2)	C12—C4—C14—C17	58.0(4)	C3—O4—C15B—C16	157.0(5)
O5—C5—C6—O6	−4.0(3)	C3—C4—C14—C17	−123.8(3)	C1—C2—C19—C20	116.0(2)
C13—C5—C6—O6	178.57(18)	C12—C4—C14—C15B	−161.4(5)	C3—C2—C19—C20	−64.7(3)
O5—C5—C6—C7	177.96(19)	C3—C4—C14—C15B	16.8(5)	C2—C19—C20—C21	−81.2(3)
O6—C6—C7—C8	179.39(19)	C12—C4—C14—C15A	163.9(3)	C19—C20—C21—C23	−176.2(2)
C13—C9—C10—O2	−177.14(18)	C3—C4—C14—C15A	−17.9(3)	C19—C20—C21—C22	4.0(4)

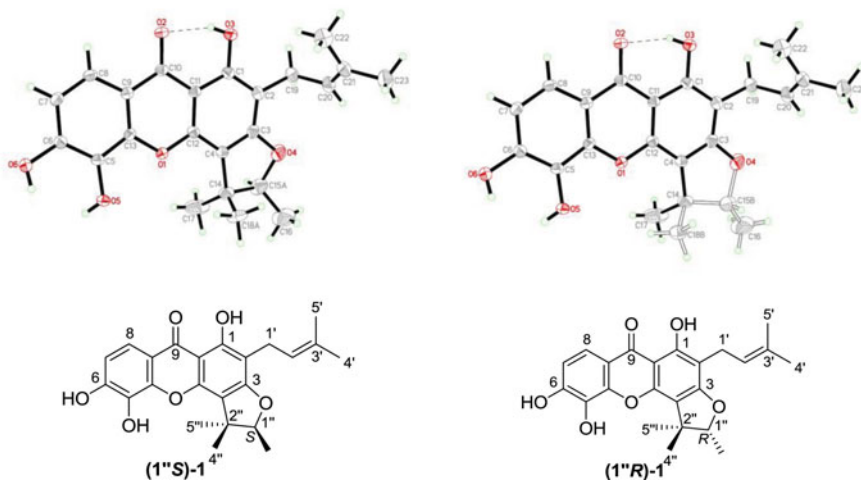


**Figure 2.** Molecular structure of **1**, showing displacement ellipsoids drawn at 30% probability level and the atom-labeling scheme. Open bond is a minor component. Intramolecular O—H...O hydrogen bond was drawn as a dashed line.

From the X-ray molecular structure in Fig. 2, the three-ring system of xanthone skeleton [C<sub>1</sub>–C<sub>13</sub>/O<sub>1</sub>] is essentially planar. The O<sub>3</sub>, O<sub>5</sub>, and O<sub>6</sub> hydroxyl O atoms lie close to their attached benzene rings with deviations +0.023(2) Å, –0.031(2) Å, and –0.008(2) Å for atoms O<sub>3</sub>, O<sub>5</sub>, and O<sub>6</sub>, respectively, and the torsion angles O<sub>3</sub>–C<sub>1</sub>–C<sub>2</sub>–C<sub>3</sub> =

**(a)** major conformational component (62.8%)

**(b)** minor conformational component (37.2 %)



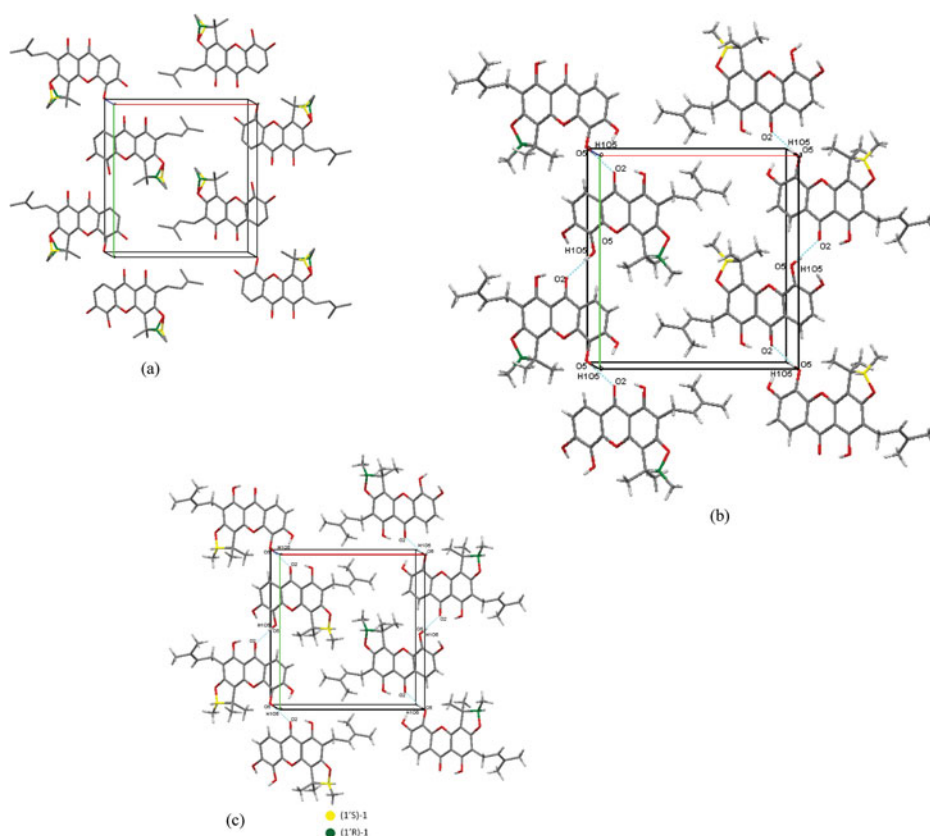
**Figure 3.** The molecular structure of (**1**) showing; **(a)** major and **(b)** minor conformational components.

**Table 3.** Hydrogen bond lengths (Å) and angles (°) for **1**

D—H...A	D—H	H...A	D...A	D—H...A
O <sub>3</sub> —H <sub>1</sub> O <sub>3</sub> ...O <sub>2</sub>	0.77(3)	1.85(3)	2.583(2)	157(3)
O <sub>5</sub> —H <sub>1</sub> O <sub>5</sub> ...O <sub>2</sub> <sup>a</sup>	0.82(4)	1.82(4)	2.636(2)	169(3)
O <sub>6</sub> —H <sub>1</sub> O <sub>6</sub> ...O <sub>5</sub> <sup>b</sup>	0.84(3)	1.99(3)	2.779(2)	158(3)
C <sub>17</sub> —H <sub>17A</sub> ...O <sub>2</sub> <sup>c</sup>	0.96	2.60	3.520(4)	161
C <sub>22</sub> —H <sub>22A</sub> ...O <sub>3</sub>	0.96	2.59	3.519(3)	163

Symmetry codes: <sup>a</sup>  $-x, -1/2 + y, 1/2 - z$ ; <sup>b</sup>  $-x, -y, 1 - z$ ; <sup>c</sup>  $x, 1/2 - y, -1/2 + z$ .

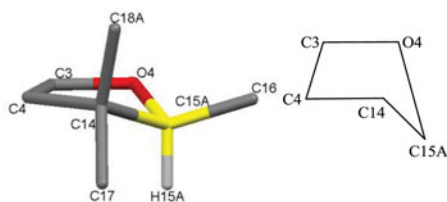
$-177.11(18)^\circ$ ,  $\text{O}_5\text{—C}_5\text{—C}_6\text{—C}_7 = 177.96(19)^\circ$ , and  $\text{O}_5\text{—C}_5\text{—C}_6\text{—C}_7 = 179.39(19)^\circ$ . The  $\alpha,\alpha,\beta$ -trimethyldihydrofuran ( $\text{C}_3\text{—C}_4/\text{C}_{14}\text{—C}_{18}/\text{O}_4$ ) moiety was found to be disordered over two sites of the major and minor conformational components with occupancy ratio of 0.628(1) and 0.372(1), respectively. It was interesting to note that this occupancy ratio corresponds to the dihydrofuran ring strain and steric hindrance between the two methyl substituents at positions 1'' and 2'' on the dihydrofuran ring.



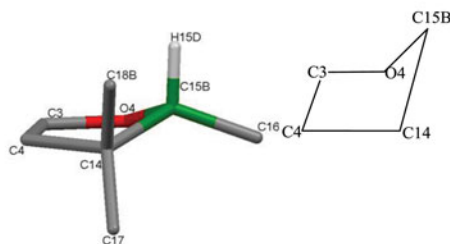
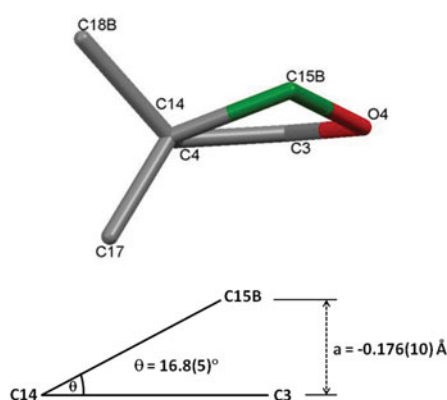
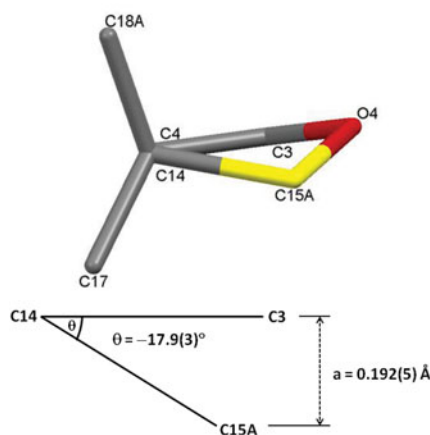
**Figure 4.** The crystal packing of **1**, showing the existence of both (1''S)-**1** and (1''R)-**1** (a) both major and minor conformational components, (b) only the major conformational component and (c) only the minor conformational component. H atoms were omitted for clarity.



(a) major conformational component

Torsion angle C3-C4-C14-C15A ( $\theta$ ) =  $-17.9(3)^\circ$ 

(b) minor conformational component

Torsion angle C3-C4-C14-C15B ( $\theta$ ) =  $16.8(5)^\circ$ 

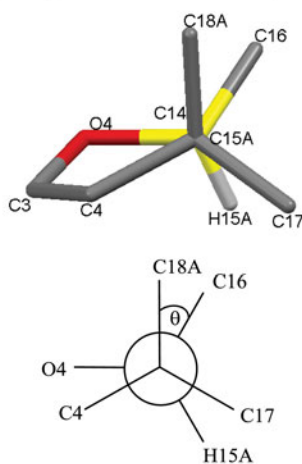
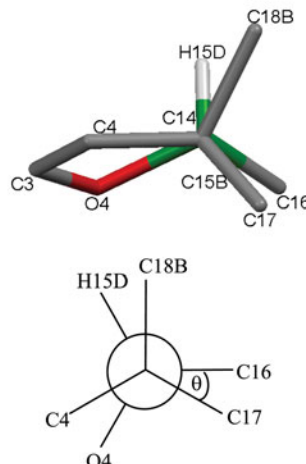
**Figure 5.** The torsion angles of C<sub>3</sub>-C<sub>4</sub>-C<sub>14</sub>-C<sub>15A/5B</sub> and the puckering values of atom C<sub>15</sub> in the (a) major and (b) minor conformational components, showing the strain of the dihydrofuran ring of **1**.

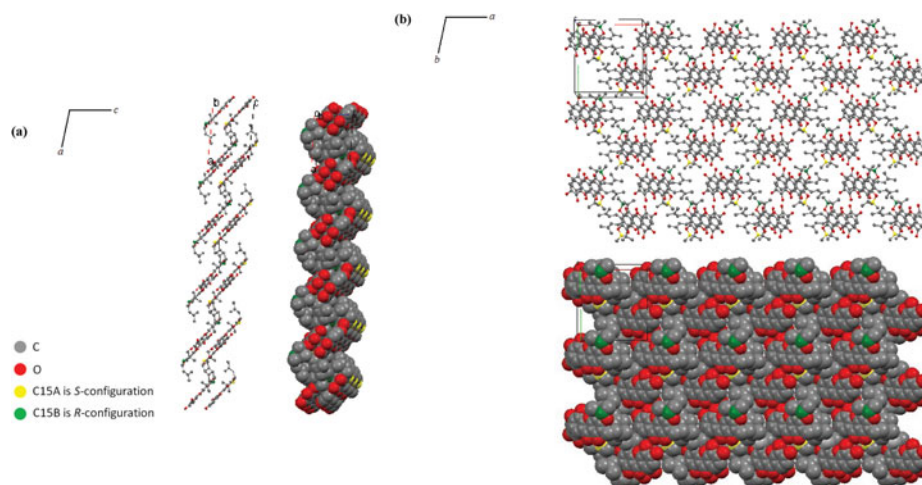
To understand the existence of the major and minor conformational components, two important criteria's in each component need to be considered which are the dihydrofuran ring strain and the steric hindrance between the two methyl groups at position 1'' and 2'' or at atoms C<sub>14</sub> and C<sub>15</sub>. For the dihydrofuran ring strain, the dihydrofuran ring of both components was in an envelope conformation with the different puckering values of atom C<sub>15</sub>, in which the major component (A), atom C<sub>15A</sub> having the deviation of 0.192(5) Å from the mean plane of C<sub>3</sub>/C<sub>4</sub>/C<sub>14</sub>/O<sub>4</sub> (see Fig. 5a) and with puckering parameters  $Q = 0.305(3)$  Å and  $\phi = 141.4(6)^\circ$  [14] and the torsion angle C<sub>3</sub>-C<sub>4</sub>-C<sub>14</sub>-C<sub>15A</sub> =  $-17.9(3)^\circ$ , whereas the minor component (B), the puckering atom C<sub>15B</sub> having the deviation of  $-0.176(10)$  Å from the mean plane of C<sub>3</sub>/C<sub>4</sub>/C<sub>14</sub>/O<sub>4</sub> (see Fig. 5b) and with puckering parameters  $Q = 0.278(5)$  Å and  $\phi = 323.3(10)^\circ$  [14] and the torsion angle C<sub>3</sub>-C<sub>4</sub>-C<sub>14</sub>-C<sub>15B</sub> =  $16.8(5)^\circ$ . It can be seen that the dihydrofuran ring with puckering value of 0.192(5) Å and the torsion angle C<sub>3</sub>-C<sub>4</sub>-C<sub>14</sub>-C<sub>15A</sub> =  $-17.9(3)^\circ$  should have less strain than that of the one with the puckering value of  $-0.176(10)$  Å and the torsion angle C<sub>3</sub>-C<sub>4</sub>-C<sub>14</sub>-C<sub>15B</sub> =  $16.8(5)^\circ$ . It can be concluded that the former component should exist as a major component, whereas the latter one as minor conformational component due to the dihydrofuran ring strain.

**Table 4.** Torsion angles ( $^\circ$ ) in the  $\alpha,\alpha,\beta$ -trimethyldihydrofuran ring of the major and minor conformational components

Major component		Minor component	
Torsion angles	( $^\circ$ )	Torsion angles	( $^\circ$ )
C <sub>18A</sub> –C <sub>14</sub> –C <sub>15A</sub> –C <sub>16</sub>	35.80	C <sub>17</sub> –C <sub>14</sub> –C <sub>15B</sub> –C <sub>16</sub>	–11.70
C <sub>17</sub> –C <sub>14</sub> –C <sub>15A</sub> –C <sub>16</sub>	–92.90	C <sub>18B</sub> –C <sub>14</sub> –C <sub>15B</sub> –C <sub>16</sub>	98.30
C <sub>17</sub> –C <sub>14</sub> –C <sub>15A</sub> –H <sub>15A</sub>	28.54	C <sub>18B</sub> –C <sub>14</sub> –C <sub>15B</sub> –H <sub>15D</sub>	–28.70
C <sub>4</sub> –C <sub>14</sub> –C <sub>15A</sub> –H <sub>15A</sub>	–87.18	C <sub>4</sub> –C <sub>14</sub> –C <sub>15B</sub> –H <sub>15D</sub>	89.11
C <sub>4</sub> –C <sub>14</sub> –C <sub>15A</sub> –O <sub>4</sub>	27.93	C <sub>4</sub> –C <sub>14</sub> –C <sub>15B</sub> –O <sub>4</sub>	–24.61
C <sub>18A</sub> –C <sub>14</sub> –C <sub>15A</sub> –O <sub>4</sub>	–87.63	C <sub>17</sub> –C <sub>14</sub> –C <sub>15B</sub> –O <sub>4</sub>	107.57

In addition, the steric hindrance between the methyl substituents on the dihydrofuran ring can be indicated by the torsion angles which were summarized in Table 4 and Fig. 6. The key is to consider the torsion angle (Fig. 6); the torsion angle C<sub>18A</sub>–C<sub>14</sub>–C<sub>15A</sub>–C<sub>16</sub> of the major component (A) was 35.80° (Table 4 and Fig. 6a), whereas the torsion angle at C<sub>17</sub>–C<sub>14</sub>–C<sub>15B</sub>–C<sub>16</sub> of the minor component (B) was –11.70° (Table 4 and Fig. 6b). The component with the wider torsion angle (35.80°) has definitely less steric hindrance than the narrow one (–11.70°). It is suggested that the former component with the wider torsion angle should exist as the major component, whereas the latter one as the minor component. In addition, to avoid steric hindrance in the molecule of **1**, the 3-methylbut-2-enyl moiety (C<sub>19</sub>–C<sub>23</sub>) is rotated by 73.10(12)° with respect to the C<sub>1</sub>–C<sub>4</sub>/C<sub>11</sub>–C<sub>12</sub>

**(a)** major conformational component**(b)** minor conformational componentTorsion angle C<sub>18A</sub>–C<sub>14</sub>–C<sub>15A</sub>–C<sub>16</sub> ( $\theta$ ) = 35.80°Torsion angle C<sub>17</sub>–C<sub>14</sub>–C<sub>15B</sub>–C<sub>16</sub> ( $\theta$ ) = –11.70°**Figure 6.** Conformations of the  $\alpha,\alpha,\beta$ -trimethyldihydrofuran ring and the substituent groups, showing the steric hindrance between the two methyl groups at positions C<sub>14</sub> and C<sub>15</sub> in the **(a)** major and **(b)** minor conformational components of **1**.



**Figure 7.** The crystal packing of **1**, H atoms were omitted for clarity; (a) showing the screw chains (b) showing the 2D network parallel to the *ab* plane.

ring. The orientation of the 3-methylbut-2-enyl moiety (C<sub>9</sub>–C<sub>23</sub>) at atom C<sub>2</sub> can be indicated by the torsion angle of C<sub>2</sub>–C<sub>19</sub>–C<sub>20</sub>–C<sub>21</sub> = –81.2(3)°, C<sub>19</sub>–C<sub>20</sub>–C<sub>21</sub>–C<sub>22</sub> = 4.01(4)°, and C<sub>19</sub>–C<sub>20</sub>–C<sub>21</sub>–C<sub>23</sub> = –176.2(2)° (Table 2). Intramolecular O<sub>2</sub>–H<sub>1</sub>O<sub>3</sub>...O<sub>2</sub> hydrogen bond (Table 3 and Fig. 2) generate S(6) ring motif [15] which help to stabilize the planarity of the xanthone ring system.

The intra- and intermolecular hydrogen bonds (Table 3) were analyzed with the program *PLATON* [12]. In the crystal structure of **1**, the molecules are linked by O<sub>5</sub>–H<sub>1</sub>O<sub>5</sub>...O<sub>2</sub> hydrogen bonds (Fig. 3 and Table 3) into screw chains (Fig. 7a). These screw chains are further linked into two dimensional network parallel to the *ab* plane (Fig. 7b) by O<sub>6</sub>–H<sub>1</sub>O<sub>6</sub>...O<sub>5</sub> hydrogen bonds (Table 3) and are stacked along the *c*-axis by  $\pi$ ... $\pi$  interactions with the centroid–centroid distances of Cg<sub>1</sub>...Cg<sub>2</sub><sup>d</sup> = 3.7520(11) Å and Cg<sub>1</sub>...Cg<sub>3</sub><sup>c</sup> = 3.7891(11) Å; Cg<sub>1</sub>, Cg<sub>2</sub>, and Cg<sub>3</sub> are the centroids of C<sub>9</sub>–C<sub>13</sub>/O<sub>1</sub>, C<sub>1</sub>–C<sub>4</sub>/C<sub>11</sub>–C<sub>12</sub> and C<sub>5</sub>–C<sub>9</sub>/C<sub>13</sub> rings, respectively, with symmetry codes: *c* = *x*, 1/2 – *y*, –1/2 + *z*, and *d* = *x*, 1/2 – *y*, 1/2 + *z*. The crystal structure is consolidated and stabilized by intra- and intermolecular O–H...O hydrogen bonds together with C–H...O interactions. Details of these interactions bond lengths and bond angles are listed in Table 3. The importance of hydrogen bonds in natural products was well documented as it was known that the numbers and positions of hydrogen donors are important and related to their bioactivities. Compound **1** which is a natural product bearing a rich source of hydrogen bond donors (CH and OH) and acceptors (O) and possesses potent nitric oxide inhibitory activity and antibacterial activities [9]. Hydrogen bonds are essential to maintain the bioactive conformation and for recognition of the binding sites of the corresponding receptors [16].

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## Supplementary Material

CCDC 910092 contains the supplementary crystallographic data for this article. These data can be obtained free of charge at [http://www.ccdc.cam.ac.uk/data\\_request/cif](http://www.ccdc.cam.ac.uk/data_request/cif), or by e-mailing [data\\_request@ccdc.cam.ac.uk](mailto:data_request@ccdc.cam.ac.uk), or by contacting the Cambridge Crystallographic Data Centre, 12, Union Road, Cambridge CB2 1EZ, UK; Fax: +44 1223 336033; e-mail: [deposit@ccdc.cam.ac.uk](mailto:deposit@ccdc.cam.ac.uk).

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