

3,6,9-Trimethyl-2,3-dihydrobenzo[de]-
chromene-7,8-dioneHoong-Kun Fun,^{a*}
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Key indicators

Single-crystal X-ray study

T = 100 K

Mean $\sigma(C-C)$ = 0.005 Å

R factor = 0.061

wR factor = 0.166

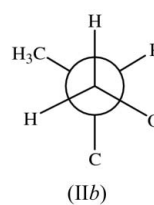
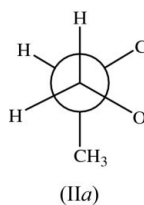
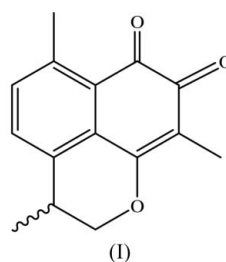
Data-to-parameter ratio = 10.8

For details of how these key indicators were
automatically derived from the article, see
<http://journals.iucr.org/e>.

The title sesquiterpene *ortho*-naphthoquinone compound, C₁₅H₁₄O₃, was isolated from *Thespesia populnea*. There are four independent molecules (*A*, *B*, *C* and *D*) in the asymmetric unit. The conformations of molecules *A* and *B* differ from those observed for *C* and *D*. In all four molecules, the pyran ring adopts an envelope conformation, with the methylene C atom as the flap atom. The molecules are arranged in layers parallel to the (100) plane, and are interconnected into a three-dimensional network by C—H···O interactions.

Comment

Thespesia populnea, or Po-ta-lae in Thai, is a plant in the *Malvaceae* with mainly tropical and subtropical worldwide distribution. The heartwood of *T. populnea* is a rich source of sesquiterpenoid quinines (Milbrodt *et al.*, 1997). The title compound, (I), also known as mansonone E (Marini Bettolo *et al.*, 1965; Tanaka *et al.*, 1966; Kim *et al.*, 1996), was isolated from the heartwood of *T. populnea* which was collected from Suratthani province in the southern part of Thailand. As part of our ongoing studies on the structure and biological activities of natural products from medicinal plants (Boonnak *et al.*, 2005; Chantrapromma *et al.*, 2005; Chantrapromma, Boonnak *et al.*, 2006; Chantrapromma, Boonsri *et al.*, 2006; Chantrapromma, Fun *et al.*, 2006; Fun *et al.*, 2005), we have undertaken the X-ray crystal structure determination of (I). The results of our biological activity investigation of (I) show strong cytotoxic activity and moderate antibacterial activity against *B. subtilis*.



Compound (I) crystallizes with four independent molecules (*A*, *B*, *C* and *D*) per asymmetric unit (Fig. 1). The dimensions

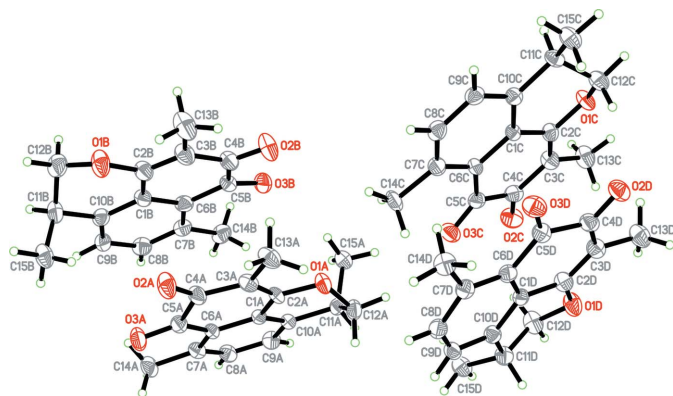


Figure 1
The asymmetric unit of the title compound, showing 50% probability displacement ellipsoids and the atomic numbering scheme.

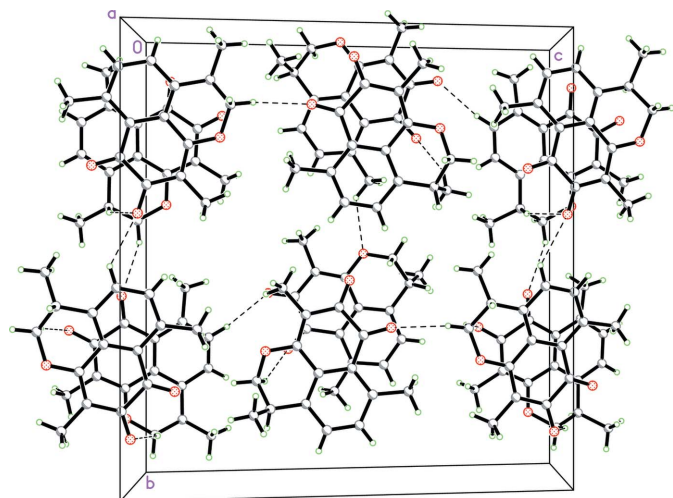


Figure 2
The crystal packing of (I), viewed approximately along the *a* axis. Hydrogen bonds are shown as dashed lines.

of all four molecules are very similar, except for slight deviations in the bond angles subtended at C11 (Table 1), with bond lengths and angles within normal ranges (Allen *et al.*, 1987). The conformations of molecules *A* and *B* are similar, the methylene and methyl groups in the pyran rings being on opposite sides of the mean plane through the molecule, while molecules *C* and *D* have similar conformations with these groups on the same side of the plane. The projections of the groups attached at C11 and C12, viewed along the C12–C11 bond, are shown in (IIa) for molecules *A* and *B*, and (IIb) for *C* and *D*.

The naphthoquinone ring system (C1–C10) is essentially planar, with atom C4 deviating by a maximum of 0.090 (4), 0.101 (4), 0.073 (4) and 0.075 (4) Å for molecules *A*, *B*, *C* and *D*, respectively. The pyran ring adopts an envelope conformations in all four molecules, with atom C12 displaced from the C1/C2/C10/C11/O1 plane by 0.640 (5), 0.640 (7) 0.636 (5) and 0.596 (6) Å for molecules *A*, *B*, *C* and *D*, respectively. The puckering parameters (Cremer & Pople, 1975) are $Q = 0.468$ (4) Å, $\theta = 57.2$ (5) and $\varphi = 293.8$ (5)° for molecule *A*, $Q =$

0.469 (5) Å, $\theta = 56.9$ (6) and $\varphi = 292.7$ (6)° for molecule *B*, $Q = 0.465$ (4) Å, $\theta = 123.0$ (5) and $\varphi = 115.1$ (6)° for molecule *C*, and $Q = 0.434$ (4) Å, $\theta = 122.5$ (5) and $\varphi = 123.1$ (6)° for molecule *D*. In molecules *A* and *B*, the methyl group at atom C11 is axially attached, while in *C* and *D* it is equatorially attached. Molecules *A*, *B* and *C*, *D* are related by conformational enantiomerism.

In the crystal structure, the molecules are stacked in layers parallel to the (100) plane. The molecules in a layer and in adjacent layers are linked through C–H...O intermolecular hydrogen bonds (Table 2), forming a three-dimensional network (Fig. 2).

Experimental

Air-dried heartwood of *T. populnea* (2.1 kg) was extracted with CH₂Cl₂ over a period of 5 d at room temperature. The CH₂Cl₂ extract (20 ml) was evaporated under reduced pressure, yielding an orange-brown gum (37.5 g), which was subjected to silica-gel column chromatography, affording eight fractions (F1–F8). Fraction F1 was subjected to repeated column chromatography, affording the title compound, (I). Single crystals of (I) were obtained after several days by recrystallization from MeOH–CH₂Cl₂ (3:7 v/v) (m.p. 381–383 K).

Crystal data

C ₁₅ H ₁₄ O ₃	$V = 2384.45$ (6) Å ³
$M_r = 242.26$	$Z = 8$
Monoclinic, $P2_1$	Mo $K\alpha$ radiation
$a = 7.1615$ (1) Å	$\mu = 0.09$ mm ⁻¹
$b = 18.6558$ (3) Å	$T = 100.0$ (1) K
$c = 18.0323$ (3) Å	$0.32 \times 0.20 \times 0.10$ mm
$\beta = 98.216$ (1)°	

Data collection

Bruker SMART APEX2 CCD area-detector diffractometer	30798 measured reflections
Absorption correction: multi-scan (SADABS; Bruker, 2005)	7133 independent reflections
$T_{\min} = 0.971$, $T_{\max} = 0.991$	4478 reflections with $I > 2\sigma(I)$
	$R_{\text{int}} = 0.080$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.061$	1 restraint
$wR(F^2) = 0.166$	H-atom parameters constrained
$S = 1.01$	$\Delta\rho_{\text{max}} = 0.32$ e Å ⁻³
7133 reflections	$\Delta\rho_{\text{min}} = -0.23$ e Å ⁻³
661 parameters	

Table 1

Selected geometric parameters (Å, °).

O1A–C2A	1.354 (5)	O1C–C2C	1.351 (5)
O1A–C12A	1.450 (5)	O1C–C12C	1.448 (5)
O2A–C4A	1.221 (5)	O2C–C4C	1.227 (5)
O3A–C5A	1.204 (5)	O3C–C5C	1.214 (5)
O1B–C2B	1.347 (5)	O1D–C2D	1.359 (5)
O1B–C12B	1.452 (6)	O1D–C12D	1.448 (5)
O2B–C4B	1.226 (5)	O2D–C4D	1.227 (5)
O3B–C5B	1.207 (5)	O3D–C5D	1.211 (5)
C12A–C11A–C15A	112.7 (4)	C12C–C11C–C15C	108.9 (3)
C12B–C11B–C15B	113.6 (4)	C12D–C11D–C15D	108.0 (4)

Table 2
Hydrogen-bond geometry (Å, °).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
C9C—H9CA \cdots O2C ⁱ	0.93	2.45	3.185 (5)	136
C8D—H8DA \cdots O2B ⁱⁱ	0.93	2.43	3.352 (5)	169
C11D—H11D \cdots O2C ⁱⁱ	0.98	2.55	3.449 (5)	152
C12A—H12A \cdots O3B ⁱⁱⁱ	0.97	2.43	3.248 (5)	142
C12C—H12E \cdots O3A ⁱⁱⁱ	0.97	2.43	3.368 (5)	162
C12C—H12F \cdots O2D	0.97	2.43	3.349 (5)	158
C12D—H12G \cdots O3D ^{iv}	0.97	2.46	3.343 (5)	151
C12D—H12H \cdots O2C	0.97	2.51	3.358 (6)	146
C14B—H14F \cdots O1B ^v	0.96	2.56	3.273 (5)	131
C15A—H15B \cdots O3B	0.96	2.52	3.410 (5)	154

Symmetry codes: (i) $-x+1, y-\frac{1}{2}, -z+2$; (ii) $x+1, y, z$; (iii) $x, y, z+1$; (iv) $-x+2, y+\frac{1}{2}, -z+2$; (v) $-x, y+\frac{1}{2}, -z+1$.

H atoms were placed in calculated positions, with C—H = 0.93–0.98 Å. The U_{iso} values were set equal to $1.5U_{eq}$ of the carrier atom for methyl H atoms and $1.2U_{eq}$ for the remaining H atoms. A rotating-group model was used for the methyl groups. In the absence of significant anomalous scattering effects, Friedel pairs were averaged.

Data collection: *APEX2* (Bruker, 2005); cell refinement: *APEX2*; data reduction: *SAINTE* (Bruker, 2005); program(s) used to solve structure: *SHELXTL* (Sheldrick, 1998); program(s) used to refine structure: *SHELXTL*; molecular graphics: *SHELXTL*; software used to prepare material for publication: *SHELXTL* and *PLATON* (Spek, 2003).

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