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Key indicators

Single-crystal X-ray study T = 293 KMean σ (C–C) = 0.007 Å R factor = 0.045 wR factor = 0.112 Data-to-parameter ratio = 7.3

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

Obacunone

The title compound, $C_{26}H_{30}O_7$, crystallizes with three independent molecules in the asymmetric unit. The crystal structure is stabilized by intra- and intermolecular $C-H\cdots O$ hydrogen-bonding interactions.

Comment

Cortex dictamni, bai-xian-pi in Chinese, is the dry root bark of Dictamnus dasycarpus Turcz (Rutaceae). It is officially listed in the Chinese pharmacopoeia (Chinese Pharmacopoeia Commission, 1995) and widely used for the treatment of jaundice, coughs and rheumatism (Jiangsu New Medical College, 1986). A number of phytochemical studies have been made of its constituents and various compounds, including furoquinoline alkaloids (Ishii et al., 1974; Deshpande & Shastri, 1977), limonoids (Zhao, Wolfender, Hostettmann, Xu & Qin, 1998) and sesquiterpenes (Chang et al., 2001; Takeuchi et al., 1993; Zhao, Wolfender, Hostettmann, Li et al., 1998), have been reported. Within the frame of a research project aimed at finding natural products with a selective immunosuppressive activity for drug discovery, we have reported that the aqueous extract of cortex dictamni possesses an attenuating effect on both cellular and humoral immune responses in mice (Wang et al., 1992). Here we report the crystal structure obacunone, 14,15:21,23-diepoxy-4,4,8-trimethyl-A,D-diof homo-24-nor-4,17-dioxachola-1,20,22-triene-3,7,16-trione, (I), a main component of cortex dictamni, which has recently attracted much attention because of its wide variety of biological activities (Lee et al., 2005; Ruberto et al., 2002; Tanaka et al., 2001).



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The asymmetric unit of (I) consists of three crystallographically independent molecules (molecule 1: O1–O7/C1– C26; molecule 2: O8–O14/C27–C52; molecule 3: O15–O21/

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Figure 1

The structures of (a) molecule 1, (b) molecule 2 and (c) molecule 3 in the asymmetric unit of (I), showing 30% probability displacement ellipsoids and the atom-numbering scheme.

C53–C78; Fig. 1*a*, 1*b* and 1*c*). All bond lengths and angles in (I) are within normal ranges (Allen *et al.*, 1987). The A/B, B/C and C/D ring junctions are all *trans*-fused. The five-membered ring is equatorially attached to ring *D*. The seven-membered heterocyclic ring *A* adopts a twist-boat conformation (Cremer & Pople, 1975; Table 2). The six-membered ring *B* assumes a chair conformation, while rings *C* and *D* show twist-boat conformations (Table 2). The crystal packing of (I) is stabilized by an extended three-dimensional network of intra- and



Figure 2

The crystal packing of (I), viewed along the *a* axis. Intermolecular hydrogen bonds are shown as dashed lines.

intermolecular $C-H\cdots O$ hydrogen-bonding interactions (Table 1, Fig. 2).

Experimental

All the solvents and reagents were commercial and purified according to standard laboratory techniques. Powdered root bark (5.0 kg) of D. dasycarpus was extracted three times with 60% EtOH under reflux. The total filtrate was evaporated under reduced pressure to remove EtOH; the extract was suspended in distilled water and partitioned with CH₂Cl₂ and EtOAc successively, to yield CH₂Cl₂ (148 g) and EtOAc (18 g) extracts, respectively. The CH₂Cl₂ extract (100 g) was subjected to silica gel column chromatography, eluted with CH₂Cl₂/ MeOH (10:1 and 2:1), and further purified by flash column chromatography (silica gel) using a gradient of CH₂Cl₂-MeOH to afford the title compound (1.0 g). The structure of (I) was confirmed by comparison of MS, ¹H NMR, ¹³C NMR and physical data with those reported in the literature (Dreyer, 1965; Pailer, et al., 1965; Kang, et al., 1986; Zhang, et al., 2005). Single crystals of (I) suitable for X-ray crystallographic analysis were obtained by slow evaporation of an MeOH–CH₂Cl₂ (2:1 v/v) solution (yield 65%).

Crystal	data
$C_{26}H_{30}C_{26}$	D_7

 $\begin{array}{l}
 M_r = 454.50 \\
 Orthorhombic, P2_12_12_1 \\
 a = 12.531 (2) Å \\
 b = 23.041 (5) Å \\
 c = 23.649 (5) Å \\
 V = 6828 (2) Å^3 \\
 \end{array}$

Z = 12 $D_x = 1.326 \text{ Mg m}^{-3}$ Mo K α radiation $\mu = 0.10 \text{ mm}^{-1}$ T = 293 (2) K Block, colourless $0.30 \times 0.25 \times 0.20 \text{ mm}$ Data collection

Bruker SMART APEX areadetector diffractometer φ and ω scans Absorption correction: multi-scan (*SADABS*; Sheldrick, 1996) $T_{\min} = 0.972, T_{\max} = 0.981$

Refinement

Refinement on F^2 $R[F^2 > 2\sigma(F^2)] = 0.045$ $wR(F^2) = 0.113$ S = 0.846625 reflections 909 parameters H-atom parameters constrained 34845 measured reflections 6625 independent reflections 3328 reflections with $I > 2\sigma(I)$ $R_{\text{int}} = 0.098$ $\theta_{\text{max}} = 25.0^{\circ}$

$$\begin{split} w &= 1/[\sigma^2(F_o^2) + (0.0501P)^2] \\ \text{where } P &= (F_o^2 + 2F_c^2)/3 \\ (\Delta/\sigma)_{\text{max}} < 0.001 \\ \Delta\rho_{\text{max}} &= 0.20 \text{ e } \text{\AA}^{-3} \\ \Delta\rho_{\text{min}} &= -0.21 \text{ e } \text{\AA}^{-3} \\ \text{Extinction correction: SHELXL97} \\ \text{Extinction coefficient: } 0.00153 (17) \end{split}$$

Table 1

Hydrogen-bond geometry (Å, °).

$D - H \cdot \cdot \cdot A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - H \cdots A$
C18-H18A····O1	0.96	2.46	3.041 (7)	119
$C22-H22A\cdots O5$	0.96	2.55	2.970 (6)	107
C44−H44 <i>B</i> ···O8	0.96	2.49	3.033 (6)	116
C55-H55···O13	0.93	2.52	3.312 (5)	143
C61-H61···O10	0.97	2.57	3.448 (4)	149
C64-H64 <i>B</i> ···O13	0.97	2.57	3.403 (6)	144
C70−H70C···O15	0.96	2.56	2.925 (5)	103
C74−H74A···O19	0.96	2.55	2.964 (5)	106
C9−H9···O17 ⁱ	0.98	2.53	3.460 (6)	159
$C22-H22C\cdots O17^{i}$	0.96	2.58	3.439 (6)	149
C23−H23···O2 ⁱⁱ	0.93	2.48	3.188 (8)	133
$C37 - H37B \cdot \cdot \cdot O19^{i}$	0.97	2.57	3.424 (5)	147
$C37 - H37B \cdot \cdot \cdot O20^{i}$	0.97	2.58	3.299 (5)	131
$C72 - H72A \cdots O8^{iii}$	0.96	2.56	3.409 (7)	148
$C73-H73B\cdots O21^{iv}$	0.96	2.55	3.430 (7)	152

Symmetry codes: (i) -x + 1, $y - \frac{1}{2}$, $-z + \frac{3}{2}$, (ii) x + 1, y, z; (iii) $x + \frac{1}{2}$, $-y + \frac{1}{2}$, -z + 1; (iv) $-x + \frac{3}{2}$, -y + 1, $z - \frac{1}{2}$.

 Table 2

 Puckering parameters (Å, °) for seven- and six-membered rings..

	<i>q</i> 2	<i>q</i> 3	φ_2	φ_3	Q	θ_2
Molecule 1						
Ring A	0.821 (4)	0.273 (4)	-20.1(3)	75.6 (10)	0.866 (4)	71.6 (3)
Ring B	0.053 (4)	0.528 (4)	103 (5)		0.503 (4)	5.8 (4)
Ring C	0.755 (5)	-0.102(5)	-159.5(3)		0.764 (5)	97.7 (4)
Ring D	0.495 (6)	-0.190(6)	97.4 (6)		0.530 (6)	111.0 (7)
Molecule 2						
Ring A	0.816 (4)	0.301 (5)	-16.2(4)	70.8 (8)	0.870 (4)	69.8 (3)
Ring B	0.100 (4)	0.527 (4)	87 (3)		0.536 (4)	10.7 (4)
Ring C	0.778 (5)	-0.108(4)	-152.0(3)		0.786 (5)	97.9 (3)
Ring D	0.480(5)	-0.200(4)	96.6 (6)		0.520 (5)	112.7 (5)
Molecule 3						
Ring A	0.901 (4)	0.240 (4)	-36.3(3)	78.2 (10)	0.932 (4)	75.1 (2)
Ring B	0.085 (4)	0.536 (4)	76 (3)		0.542 (4)	9.0 (4)
Ring C	0.757 (4)	-0.099(4)	-154.7(3)		0.763 (4)	97.5 (3)
Ring D	0.537 (4)	-0.166 (4)	95.7 (4)		0.562 (4)	107.2 (4)

Owing to the lack of significant anomalous scattering, Friedel pairs were merged before the final refinement and the absolute configuration could not be assigned. All H atoms were placed in geometrically idealized positions and constrained to ride on their parent atoms, with C-H = 0.93–0.98 Å, and with $U_{\rm iso}$ (H) = $1.2U_{\rm eq}$ (C), or $U_{\rm iso}$ (H) = $1.5U_{\rm eq}$ (C) for methyl H atoms.

Data collection: *SMART* (Bruker, 2003); cell refinement: *SAINT* (Bruker, 2003); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997*a*); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997*a*); molecular graphics: *SHELXTL* (Sheldrick, 1997*b*); software used to prepare material for publication: *SHELXL97*.

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