

Tricoccin R6

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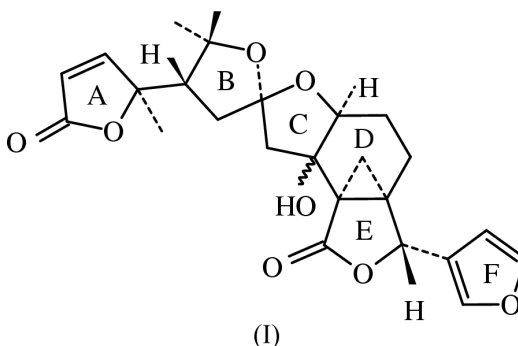
Key indicators

Single-crystal X-ray study
T = 293 K
Mean $\sigma(\text{C}-\text{C}) = 0.004 \text{ \AA}$
R factor = 0.033
wR factor = 0.092
Data-to-parameter ratio = 8.0For details of how these key indicators were automatically derived from the article, see <http://journals.iucr.org/e>.

The X-ray crystal structure of the title compound, $\text{C}_{25}\text{H}_{28}\text{O}_8$, has been determined. In the structure, both the terminal five-membered rings (*A* and *F*) are planar. The fused five-membered rings *C* and *E* are in envelope conformations, and ring *B* is in a slightly distorted half-chair conformation. The six-membered ring *D* is in a slightly distorted sofa conformation. The structure is stabilized by $\text{O}-\text{H}\cdots\text{O}$ hydrogen bonds and $\text{C}-\text{H}\cdots\text{O}$ inter- and intramolecular interactions.

Comment

The molecular structure of tricoccin R6, (I), is shown in Fig. 1. The bond geometry conforms to expectations. The puckering parameters evaluated using *PARST97* (Nardelli, 1995) show that (i) both the terminal five-membered rings (*A* and *F*) are planar, (ii) the fused five-membered ring *B* is in a slightly distorted half-chair conformation, and rings *C* and *E* are in envelope conformations [$q_2 = 0.334(2) \text{ \AA}$, $\varphi_2 = -119.7(4)^\circ$ for ring *B*, $q_2 = 0.440(2) \text{ \AA}$, $\varphi_2 = -134.7(3)^\circ$ for ring *C*, and $q_2 = 0.129(2) \text{ \AA}$, $\varphi_2 = 151.2(11)^\circ$ for ring *E*]. The six-membered ring *D* has a slightly distorted sofa conformation [$q_2 = 0.369(3)$, $q_3 = 0.364(3)$, $Q_T = 0.519(2) \text{ \AA}$, $\varphi_2 = 63.0(4)^\circ$]. Fig. 2 shows the packing diagram of the molecules. The structure is stabilized by intermolecular $\text{O}-\text{H}\cdots\text{O}$ hydrogen bonds and $\text{C}-\text{H}\cdots\text{O}$ inter- and intramolecular interactions (see Table 1).



Experimental

The title compound was isolated from *Cneorum tricoccin* L., a shrub native to coastal areas of the western Mediterranean with hairless leaves, yellow blossoms and red fruits (Herz *et al.*, 1983). It was crystallized from ethanol/acetone.

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Crystal data

$C_{25}H_{28}O_8$
 $M_r = 456.47$
 Orthorhombic, $P2_12_12_1$
 $a = 7.224$ (2) Å
 $b = 15.521$ (2) Å
 $c = 20.404$ (1) Å
 $V = 2287.8$ (7) Å³
 $Z = 4$
 $D_x = 1.325$ Mg m⁻³

Data collection

Enraf–Nonius CAD-4 diffractometer
 $\omega/2\theta$ scans
 2417 measured reflections
 2417 independent reflections
 2171 reflections with $I > 2\sigma(I)$
 $\theta_{\max} = 69.9^\circ$

Refinement

Refinement on F^2
 $R[F^2 > 2\sigma(F^2)] = 0.033$
 $wR(F^2) = 0.092$
 $S = 1.04$
 2417 reflections
 303 parameters
 H-atom parameters constrained
 $w = 1/[\sigma^2(F_o^2) + (0.0539P)^2 + 0.3026P]$
 where $P = (F_o^2 + 2F_c^2)/3$

Cu $K\alpha$ radiation
 Cell parameters from 25 reflections
 $\theta = 20\text{--}30^\circ$
 $\mu = 0.82$ mm⁻¹
 $T = 293$ (2) K
 Section from needle, colourless
 $0.25 \times 0.25 \times 0.20$ mm

$h = 0 \rightarrow 8$
 $k = 0 \rightarrow 18$
 $l = 0 \rightarrow 24$
 3 standard reflections
 frequency: 120 min
 intensity decay: <1%

$(\Delta/\sigma)_{\max} = 0.001$
 $\Delta\rho_{\max} = 0.19$ e Å⁻³
 $\Delta\rho_{\min} = -0.17$ e Å⁻³
 Extinction correction: *SHELXL97*
 Extinction coefficient: 0.0078 (4)
 Absolute structure: Flack (1983)
 Flack parameter = 0.2 (2)

Table 1

Hydrogen-bonding geometry (Å, °).

$D-H \cdots A$	$D-H$	$H \cdots A$	$D \cdots A$	$D-H \cdots A$
$C28-H28C \cdots O32$	0.96	2.41	2.894 (4)	111
$O26-H26 \cdots O25^i$	0.82	2.00	2.805 (2)	169
$C23-H23 \cdots O31^{ii}$	0.93	2.43	3.283 (3)	152

Symmetry codes: (i) $x - \frac{1}{2}, \frac{1}{2} - y, 2 - z$; (ii) $2 - x, y - \frac{1}{2}, \frac{5}{2} - z$.

The data set contains no Friedel pairs.

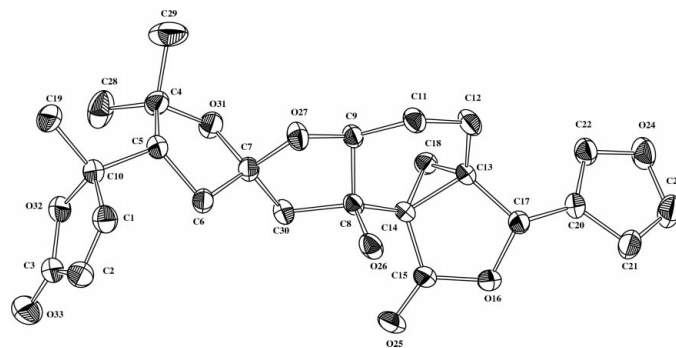


Figure 1

The molecular structure of the title compound with 30% probability displacement ellipsoids.

Data collection: *CAD-4 Software* (Enraf–Nonius, 1989); cell refinement: *SDP* (Frenz, 1978); data reduction: *CAD-4 Software*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *ORTEPIII* (Johnson & Burnett, 1998); software used to prepare material for publication: *SHELXL97* and *PARST97* (Nardelli, 1995).

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