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Key indicators

Single-crystal X-ray study T = 295 KMean σ (C–C) = 0.006 Å R factor = 0.065 wR factor = 0.179 Data-to-parameter ratio = 8.5

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

10-Acetonyl-10-hydroxy-1,6-dimethyl-1,2,10,11-tetrahydrophenanthro[1,2-b]furan-11-one

In the crystal structure of the title compound, $C_{21}H_{20}O_4$, an intramolecular O-H···O hydrogen bond is found between a hydroxyl H atom and a carbonyl O atom.

Comment

The title compound (danshenol A), (I), was extracted from Salvia yunnanensis (Xu et al., 2006) and recrystallized from ethanol. As it shows strong aldose reductase (AR) inhibitory activity (Tezuka et al., 1997) we have determined its structure.

The molecular structure of (I) is shown in Fig. 1. All four rings are coplanar and an intramolecular O-H···O hydrogen bond is found between a hydroxyl H atom and a carbonyl O atom (Fig. 2).

Experimental

The dried and powdered roots of S. yunnanensis were extracted three times over 24 h with Me₂CO at room temperature and afterwards the

Figure 1

The molecular structure of (I), showing the atom-labeling scheme. Displacement ellipsoids are drawn at the 30% probability level.

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organic papers

solvent was removed *in vacuo*. The residue was subjected to column chromatography over DM-130 porous resin and eluted with MeOH- H_2O (1:1) and 90% MeOH- H_2O (9:1). The residue of the MeOH- H_2O (9:1) fraction was partitioned between H_2O and EtOAc. The EtOAc part was subjected to silica-gel column chromatography. Mixtures of petroleum ether/EtOAc (1:0, 9:1, 8:2, 7:3, 6:4, 5:5 and 0:1) of increasing polarity were used as eluants. Seven fractions were collected and combined by monitoring with thin-layer chromatography. In this procedure, a mixture of danshenol A and danshenol C was obtained from the third fraction by silica-gel column chromatography using petroleum ether-CHCl₃–EtOAc (70/25/5) as eluants. Both compounds were separated by semi-preparative high-performance liquid chromatography using 85% MeOH-H₂O as eluant. Crystals of Danshenol A suitable for data collection were obtained by slow evaporation of an ethanol solution over a period of two weeks.

Z = 2

 $D_x = 1.322 \text{ Mg m}^{-3}$

Mo $K\alpha$ radiation

Column, colorless

 $0.30 \times 0.15 \times 0.10 \text{ mm}$

1930 independent reflections

1894 reflections with $I > 2\sigma(I)$

 $\mu = 0.09 \text{ mm}^{-1}$

T = 295 (2) K

 $R_{\rm int} = 0.048$

 $\theta_{\rm max} = 27.2^\circ$

Crystal data

 $C_{21}H_{20}O_4$ $M_r = 336.37$ Monoclinic, $P2_1$ a = 7.5180 (15) Å b = 6.7150 (13) Å c = 17.161 (3) Å $\beta = 102.75 (3)^{\circ}$ $V = 845.0 (3) \text{ Å}^{3}$

Data collection

MAC DIP 2030K diffractometer ω scans Absorption correction: none 7044 measured reflections

Refinement

Refinement on F^2	$w = 1/[\sigma^2(F_o^2) + (0.1047P)^2]$
$R[F^2 > 2\sigma(F^2)] = 0.065$	+ 0.3521P]
$wR(F^2) = 0.179$	where $P = (F_0^2 + 2F_c^2)/3$
S = 1.09	$(\Delta/\sigma)_{\rm max} = 0.002$
1930 reflections	$\Delta \rho_{\rm max} = 0.28 \text{ e} \text{ \AA}^{-3}$
227 parameters	$\Delta \rho_{\rm min} = -0.22 \text{ e } \text{\AA}^{-3}$
H-atom parameters constrained	Extinction correction: SHELXL97
	Extinction coefficient: 0.081 (15)

Table 1

Hydrogen-bond	geometry	(Å, °).	
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$D - H \cdots A$	D-H	$H \cdot \cdot \cdot A$	$D \cdot \cdot \cdot A$	$D - \mathbf{H} \cdot \cdot \cdot A$
O2−H2A···O3	0.82	2.24	2.969 (6)	148

The methyl H atoms were placed in calculated positions, with C– H = 0.96 Å, allowed to rotate but not tip and were refined using a





The molecular packing of the title compound, viewed along the b axis. Hydrogen bonds are shown as dashed lines.

riding model, with $U_{iso}(H) = 1.5U_{eq}(C)$. All other C–H H atoms were placed in geometrically idealized positions with C–H = 0.92–0.98 Å and were refined using a riding model with $U_{iso}(H) = 1.2U_{eq}(C)$. The O–H H atom was placed in an ideal position such that the O–H vector points in the direction of the nearest acceptor atom and afterwards it was refined using a riding model with $U_{iso}(H) = 1.5U_{eq}(O)$. In the absence of significant anomalous cattering effects, Friedel pairs were averaged.

Data collection: *DENZO* (Otwinowski & Minor, 1997); cell refinement: *SCALEPACK* (Otwinowski & Minor, 1997); data reduction: *SCALEPACK*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *ORTEPII* (Johnson, 1976) and *PLATON* (Spek, 2003); software used to prepare material for publication: *SHELXL97*.

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