Acta Crystallographica Section E **Structure Reports** Online

ISSN 1600-5368

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Key indicators

Single-crystal X-ray study T = 120 KMean σ (C–C) = 0.002 Å R factor = 0.050 wR factor = 0.137 Data-to-parameter ratio = 22.3

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

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A new polymorph of atranorin, a lichen paradepside

In the title compound, methyl 4-(3-formyl-2,4-dihydroxy-6methyl-benzoyloxy)-2-hydroxy-3,6-dimethylbenzoate, C₁₉H₁₈- O_8 , there are three intramolecular $O-H \cdots O$ hydrogen bonds, with lengths 2.5515 (14), 2.5711 (15), and 2.5437 (13) Å. The two aromatic rings form a dihedral angle of 60.38 (3)°, differing from that in the previously reported Pbca polymorph, viz. 84 (1)°.

Received 27 June 2002 Accepted 4 July 2002 Online 19 July 2002

Comment

The title compound, (I), was isolated from the lichen Cladonia prostrata, which was being studied for allelopathic activity in the Florida Scrub (Robbs, 1997). Atranorin has been reported to show phytotoxic activity against watercress and to stimulate growth of oats (Huneck & Schreber, 1972), and also to stimulate the growth of Rudbeckia (Robbs, 1997). The crystal structure has been reported in space group Pbca (Brassy et al., 1982), but recrystallization from ethyl acetate yielded a monoclinic polymorph, the structure of which is reported here.



The conformation of the molecule is described by the five torsion angles in Table 1, and is such that the two aromatic rings form a dihedral angle of $60.38 (3)^{\circ}$. This conformation differs from that observed in the Pbca polymorph, mainly in the torsion angle C11-O4-C1-C2, which has a value of $89 (1)^{\circ}$, causing the dihedral angle between the aromatic rings to be much more nearly orthogonal, 84 (1)°. Smaller differences in torsion angles also exist between the two polymorphs about C4–C9, 10 (1)°, and C11–C12, 8 (1)°.

The molecule in the reported polymorph exhibits three intramolecular hydrogen bonds (Table 2), one of which is bifurcated, having an intermolecular component which forms dimers about an inversion center (Fig. 2). The hydrogen bonding in the Pbca polymorph is similar. In that structure, the three intramolecular hydrogen bonds also exist, and atom O7 also has an intermolecular component. However, it does not form dimers as does the present structure, but instead forms chains, with the intermolecular acceptor being O3.

The cell dimensions at 293 K are a = 10.922 (2), b =11.257 (2), and c = 14.875 (3) Å, and $\beta = 109.60$ (2)°.



Figure 1

The atom-numbering scheme for (I), with ellipsoids at the 50% probability level, showing the intramolecular hydrogen bonds.

Experimental

The title compound was isolated from the CH₂Cl₂ extract of a sample of the lichen Cladonia prostrata, collected in Sebring, Florida. Crystals were grown from ethyl acetate.

Crystal data

$C_{19}H_{18}O_8$	$D_x = 1.484 \text{ Mg m}^{-3}$
$M_r = 374.33$	Mo $K\alpha$ radiation
Monoclinic, $P2_1/n$	Cell parameters from 5466
a = 10.929 (3) Å	reflections
b = 10.976 (3) Å	$\theta = 2.5 - 32.0^{\circ}$
c = 14.843 (3) Å	$\mu = 0.12 \text{ mm}^{-1}$
$\beta = 109.745 \ (12)^{\circ}$	T = 120 K
V = 1675.7 (7) Å ³	Prism, colorless
Z = 4	$0.40 \times 0.30 \times 0.15 \text{ mm}$

Data collection

reflections with $I > 2\sigma(I)$
= 0.025
= 32.0°
$-16 \rightarrow 16$
$-16 \rightarrow 15$
$-22 \rightarrow 22$



Figure 2

The hydrogen-bonded dimer about the the center at (1/2, 0, 1/2). Only the hydroxy H atoms are shown.

Refinement

Refinement on F^2	$w = 1/[\sigma^2(F_o^2) + (0.0551P)^2]$
$R[F^2 > 2\sigma(F^2)] = 0.050$	+ 0.7055P]
$wR(F^2) = 0.137$	where $P = (F_o^2 + 2F_c^2)/3$
S = 1.06	$(\Delta/\sigma)_{\rm max} < 0.001$
5796 reflections	$\Delta \rho_{\rm max} = 0.43 \ {\rm e} \ {\rm \AA}^{-3}$
260 parameters	$\Delta \rho_{\rm min} = -0.29 \text{ e } \text{\AA}^{-3}$
H atoms treated by a mixture of	
independent and constrained	
refinement	

Table 1

Selected geometric	parameters	(Å,	°).
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C11-O4-C1-C2	63.23 (15)	O5-C11-C12-C13	-4.84 (19)
C3-C4-C9-O1	-10.13(18)	C15-C14-C19-O8	-2.3(2)
C1-O4-C11-C12	-172.30 (11)		

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

$D - H \cdot \cdot \cdot A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - H \cdots A$
O2−H2O···O3	0.89 (2)	1.72 (2)	2.5515 (14)	152.5 (18)
O6−H6O···O8	0.90(2)	1.74 (2)	2.5711 (15)	151.1 (18)
O7−H7O···O5	0.87(2)	1.74 (2)	2.5437 (13)	151.9 (18)
$O7-H7O\cdots O5^i$	0.87 (2)	2.437 (19)	3.0251 (14)	125.1 (16)

Symmetry code: (i) 1 - x, -y, 1 - z.

H atoms on C atoms were placed in calculated positions, with C-H distances in the range 0.95–1.00 Å and thereafter treated as riding. A torsional parameter was refined for each methyl group. Hydroxy H atoms were placed from difference maps, and their coordinates were refined. $U_{iso} = 1.2U_{eq}$ of the attached atom (1.5 U_{eq} for methyl and OH groups).

Data collection: COLLECT (Nonius, 1999); cell refinement: HKL SCALEPACK (Otwinowski & Minor, 1997); data reduction: HKL DENZO (Otwinowski & Minor, 1997) and SCALEPACK; program(s) used to solve structure: SIR97 (Altomare et al., 1999); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: ORTEP-3 for Windows (Farrugia, 1997); software used to prepare material for publication: SHELXL97.

The purchase of the diffractometer was made possible by grant No. LEQSF(1999-2000)-ESH-TR-13, administered by the Louisiana Board of Regents.

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