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## Key indicators

Single-crystal X-ray study
$T=120 \mathrm{~K}$
Mean $\sigma(\mathrm{C}-\mathrm{C})=0.002 \AA$
$R$ factor $=0.050$
$\omega R$ 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.iucr.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-6-methyl-benzoyloxy)-2-hydroxy-3,6-dimethylbenzoate, $\mathrm{C}_{19} \mathrm{H}_{18}{ }^{-}$ $\mathrm{O}_{8}$, there are three intramolecular $\mathrm{O}-\mathrm{H} \cdots \mathrm{O}$ hydrogen bonds, with lengths 2.5515 (14), 2.5711 (15), and 2.5437 (13) $\AA$. The two aromatic rings form a dihedral angle of $60.38(3)^{\circ}$, differing from that in the previously reported $P b c a$ polymorph, viz. $84(1)^{\circ}$.

## 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.

(I)

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 $\mathrm{C} 11-\mathrm{O} 4-\mathrm{C} 1-\mathrm{C} 2$, which has a value of $89(1)^{\circ}$, causing the dihedral angle between the aromatic rings to be much more nearly orthogonal, $84(1)^{\circ}$. Smaller differences in torsion angles also exist between the two polymorphs about $\mathrm{C} 4-\mathrm{C} 9,10(1)^{\circ}$, and $\mathrm{C} 11-\mathrm{C} 12,8(1)^{\circ}$.

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) $\AA$, and $\beta=109.60$ (2).

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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 $\mathrm{CH}_{2} \mathrm{Cl}_{2}$ extract of a sample of the lichen Cladonia prostrata, collected in Sebring, Florida. Crystals were grown from ethyl acetate.

## Crystal data

$\mathrm{C}_{19} \mathrm{H}_{18} \mathrm{O}_{8}$
$M_{r}=374.33$
Monoclinic, $P 2_{\mathrm{d}} / n$
$a=10.929$ (3) А
$b=10.976$ (3) $\AA$
$c=14.843$ (3) A
$\beta=109.745$ (12) ${ }^{\circ}$
$V=1675.7$ (7) $\mathrm{A}^{3}$
$Z=4$

$$
\begin{aligned}
& D_{x}=1.484 \mathrm{Mg} \mathrm{~m}^{-3} \\
& \text { Mo } K \alpha \text { radiation }
\end{aligned}
$$

Cell parameters from 5466
reflections
$\theta=2.5-32.0^{\circ}$
$\mu=0.12 \mathrm{~mm}^{-1}$
$T=120 \mathrm{~K}$
Prism, colorless
$0.40 \times 0.30 \times 0.15 \mathrm{~mm}$

## Data collection

KappaCCD diffractometer with an Oxford Cryosystems Cryostream cooler

$$
\begin{aligned}
& 4257 \text { reflections with } I>2 \sigma(I) \\
& R_{\text {int }}=0.025 \\
& \theta_{\max }=32.0^{\circ} \\
& h=-16 \rightarrow 16 \\
& k=-16 \rightarrow 15 \\
& l=-22 \rightarrow 22
\end{aligned}
$$

$\omega$ scans with $\kappa$ offsets
Absorption correction: none
21781 measured reflections
5796 independent reflections


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 $F^{2}$

$$
\begin{aligned}
& w= 1 /\left[\sigma^{2}\left(F_{o}^{2}\right)+(0.0551 P)^{2}\right. \\
&+0.7055 P] \\
& \text { where } P=\left(F_{o}^{2}+2 F_{c}^{2}\right) / 3
\end{aligned}
$$

$w R\left(F^{2}\right)=0.137$
$S=1.06$
5796 reflections
260 parameters
H atoms treated by a mixture of independent and constrained refinement

Table 1
Selected geometric parameters $\left(\AA,^{\circ}\right)$.

| $\mathrm{C} 11-\mathrm{O} 4-\mathrm{C} 1-\mathrm{C} 2$ | $63.23(15)$ | $\mathrm{O} 5-\mathrm{C} 11-\mathrm{C} 12-\mathrm{C} 13$ | $-4.84(19)$ |
| :--- | ---: | ---: | :--- |
| $\mathrm{C} 3-\mathrm{C} 4-\mathrm{C} 9-\mathrm{O} 1$ | $-10.13(18)$ | $\mathrm{C} 15-\mathrm{C} 14-\mathrm{C} 19-\mathrm{O} 8$ | $-2.3(2)$ |
| $\mathrm{C} 1-\mathrm{O} 4-\mathrm{C} 11-\mathrm{C} 12$ | $-172.30(11)$ |  |  |

Table 2
Hydrogen-bonding geometry $\left(\AA^{\circ},^{\circ}\right)$.

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

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

H atoms on C atoms were placed in calculated positions, with $\mathrm{C}-$ H distances in the range $0.95-1.00 \AA$ 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. $U_{\text {iso }}=1.2 U_{\text {eq }}$ of the attached atom $\left(1.5 U_{\text {eq }}\right.$ 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.

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