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## Structure Reports

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## Key indicators

Single-crystal X-ray study
$T=293 \mathrm{~K}$
Mean $\sigma(\mathrm{C}-\mathrm{C})=0.003 \AA$
$R$ factor $=0.041$
$w R$ factor $=0.122$
Data-to-parameter ratio $=15.5$

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

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## 1-(2,6-Dihydroxyphenyl)ethanone

The title compound, $\mathrm{C}_{8} \mathrm{H}_{8} \mathrm{O}_{3}$, has several intra- and intermolecular hydrogen bonds in its crystal structure. There are two molecules in the asymmetric unit, and they are extended into infinite chains along [011] by intermolecular $\mathrm{O}-\mathrm{H} \cdots \mathrm{O}$ hydrogen bonds.

## Comment

The title compound, (I), was isolated from the extracts of cultures of the estuarine fungus (No. 3920). This substance was previously isolated from the extracts of cultures of $D$. Concentrica strain 26 A1 (Allport \& Bu'Lock, 1960). The structure of (I) was previously elucidated on the basis of spectroscopic analysis. We report here the crystal structure of (I).

(I)

The X-ray crystallographic study of (I) confirms the previously proposed molecular structure based on spectroscopic data. There are two crystallographically independent molecules in the asymmetric unit (Fig. 1). The $\mathrm{C}-\mathrm{O}$ and $\mathrm{C}-\mathrm{C}$ distances show no remarkable features (Table 1). A feature of the structure of (I) is the presence of both intra- and intermolecular $\mathrm{O}-\mathrm{H} \cdots \mathrm{O}$ hydrogen bonds between the hydroxy groups and the carbonyl O atom (Table 2), resulting in infinite chains along [011] (Fig. 2).

## Experimental

A strain of fungus (No. 3920) was isolated from an endophyte NP 159/ Morphology Type10 from Kandelia Bark Mai Po, Hong Kong, and deposited in the Department of Applied Chemistry, ZhongShan University, Guangzhou, People's Republic of China. Culture conditions: GYT medium (glucose $10 \mathrm{~g}^{-1}$, peptone $2 \mathrm{~g}^{-1}$, yeast extract $1 \mathrm{~g} \mathrm{l}^{-1}, \mathrm{NaCl} 2 \mathrm{~g}^{-1}$ ) and incubation at 298 K for 28 d . Extraction and separation of metabolite: the cultures ( 100 l ) were filtered through cheesecloth. The filtrate was concentrated to 51 below 333 K , then extracted three times by shaking with an equal volume of ethyl acetate. The extract was evaporated under reduced pressure. The combined organic extracts were subjected to silica-gel column chromatography, eluting with petroleum ether/ethyl acetate, to yield the

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title compound, (I). The compound's identity was confirmed by the NMR spectra. Crystals of (I) were obtained by evaporation of a methanol solution. ${ }^{1} \mathrm{H}$ NMR ( 300 MHz , actone- $d_{6}$ ): $\delta 2.69(s, H), 6.40$ $(d, J=8.1 \mathrm{~Hz}, \mathrm{H} 3, \mathrm{H} 5), 7.23(t, J=8.1,16.2 \mathrm{~Hz}, \mathrm{H} 4), 11.44(s, 2-\mathrm{OH}, 6-$ $\mathrm{OH})$.

## Crystal data

$\mathrm{C}_{8} \mathrm{H}_{8} \mathrm{O}_{3}$
$M_{r}=152.14$
Triclinic, $P \overline{1}$
$a=7.646$ (3) A
$b=8.325$ (3) $\AA$
$c=12.803(5) \AA$
$\alpha=73.321(6)^{\circ}$
$\beta=79.042$ (7) ${ }^{\circ}$
$\gamma=68.230(6)^{\circ}$
$V=721.8(5) \AA^{3}$
Data collection
Bruker SMART CCD area-detector diffractometer
$\varphi$ and $\omega$ scans
Absorption correction: multi-scan (SADABS; Sheldrick, 1996)
$T_{\min }=0.950, T_{\max }=0.963$
6117 measured reflections

## Refinement

Refinement on $F^{2}$
$R\left[F^{2}>2 \sigma\left(F^{2}\right)\right]=0.041$
$w R\left(F^{2}\right)=0.122$
$S=1.03$
3115 reflections
201 parameters
H-atom parameters constrained

$$
\begin{aligned}
& Z=4 \\
& D_{x}=1.400 \mathrm{Mg} \mathrm{~m}^{-3}
\end{aligned}
$$

Mo $K \alpha$ radiation
Cell parameters from 816
reflections
$\theta=2.8-26.0^{\circ}$
$\mu=0.11 \mathrm{~mm}^{-1}$
$T=293$ (2) K
Block, colorless
$0.48 \times 0.42 \times 0.35 \mathrm{~mm}$

3115 independent reflections
2065 reflections with $I>2 \sigma(I)$
$R_{\text {int }}=0.015$
$\theta_{\text {max }}=27.1^{\circ}$
$h=-9 \rightarrow 9$
$k=-10 \rightarrow 10$
$l=-15 \rightarrow 16$

$$
\begin{aligned}
w= & 1 /\left[\sigma^{2}\left(F_{o}^{2}\right)+(0.0579 P)^{2}\right. \\
& +0.1147 P]
\end{aligned}
$$

where $P=\left(F_{o}{ }^{2}+2 F_{c}{ }^{2}\right) / 3$
$(\Delta / \sigma)_{\max }<0.001$
$\Delta \rho_{\text {max }}=0.24 \mathrm{e}^{\text {A. }}{ }^{-3}$
$\Delta \rho_{\min }=-0.13 \mathrm{e}^{-3}$

Table 1
Selected bond lengths $(\AA)$.

| C1-C2 | $1.414(2)$ | C9-C14 | $1.413(2)$ |
| :--- | :--- | :--- | :--- |
| C1-C6 | $1.415(2)$ | C9-C10 | $1.420(2)$ |
| C1-C7 | $1.472(2)$ | C9-C15 | $1.461(2)$ |
| C2-O1 | $1.3519(19)$ | C10-O4 | $1.3469(19)$ |
| C2-C3 | $1.382(2)$ | C10-C11 | $1.379(2)$ |
| C3-C4 | $1.369(2)$ | C11-C12 | $1.365(3)$ |
| C4-C5 | $1.379(2)$ | C12-C13 | $1.372(3)$ |
| C5-C6 | $1.379(2)$ | C13-C14 | $1.381(2)$ |
| C6-O2 | $1.3539(18)$ | C14-O5 | $1.352(2)$ |
| C7-O3 | $1.2378(19)$ | C15-O6 | $1.2425(19)$ |
| C7-C8 | $1.489(2)$ | C15-C16 | $1.486(3)$ |

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

| $D-\mathrm{H} \cdots A$ | $D-\mathrm{H}$ | $\mathrm{H} \cdots A$ | $D \cdots A$ | $D-\mathrm{H} \cdots A$ |
| :---: | :---: | :---: | :---: | :---: |
| $\mathrm{O} 1-\mathrm{H} 1 A \cdots \mathrm{O}^{\text {i }}$ | 0.82 | 1.95 | 2.767 (2) | 176 |
| $\mathrm{O} 2-\mathrm{H} 2 A \cdots \mathrm{O} 3$ | 0.82 | 1.74 | 2.472 (2) | 148 |
| $\mathrm{O} 4-\mathrm{H} 4 A \cdots \mathrm{O} 6$ | 0.82 | 1.79 | 2.513 (2) | 146 |
| $\mathrm{O} 5-\mathrm{H} 5 A \cdots \mathrm{O} 2^{\text {ii }}$ | 0.82 | 1.97 | 2.7878 (19) | 180 |

Symmetry codes: (i) $x-1,1+y, z$; (ii) $1+x, y, z-1$.



Figure 1
The asymmetric unit of (I), showing $30 \%$ probability displacement ellipsoids and the atom-numbering scheme.


Figure 2
The packing of (I), viewed down the $b$ axis. Hydrogen bonds are shown as dashed lines.

The H atoms were positioned geometrically and were treated as riding on their parent C and O atoms, with $\mathrm{C}-\mathrm{H}$ distances in the range $0.93-0.96 \AA$ and $\mathrm{O}-\mathrm{H}$ distances of $0.82 \AA$.

Data collection: SMART (Bruker, 1999); cell refinement: SAINTPlus (Bruker, 1999; data reduction: SAINT-Plus; program(s) used to solve structure: SHELXS97 (Sheldrick, 1990); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: SHELXTL (Bruker, 1999); software used to prepare material for publication: SHELXTL.

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