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## Key indicators

Single-crystal X-ray study
$T=293 \mathrm{~K}$
Mean $\sigma(\mathrm{C}-\mathrm{C})=0.004 \AA$
$R$ factor $=0.038$
$w R$ factor $=0.091$
Data-to-parameter ratio $=9.2$

For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e.
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## 3-(a-Hydroxy-2-methoxylbenzylidene)-1-isopropylpyrrolidine-2,4-dione

The title compound, $\mathrm{C}_{15} \mathrm{H}_{17} \mathrm{NO}_{4}$, is a potent new herbicide containing the pyrrolidine-2,4-dione ring system. In the crystalline state, the molecular skeleton contains one enol grouping, which is intramolecularly hydrogen bonded to a neighboring keto O atom.

## Comment

Many compounds containing the 3-acylpyrrolidine-2,4-dione moiety are novel heterocyclic compounds with antibiotic activity; these include tenuazonic acid (Stickings, 1959), streptolydigin (Rinehart et al., 1963), tirandamycin (Rinehart et al., 1971), malonomycin (Bann et al., 1978), $\alpha$-cyclopiazonic acid (Stickings, 1959; van Rooyen, 1992) and $\beta$-cyclopiazonic acid (Holzapfel et al., 1970). All these compounds possess a 3-acyltetramic acid grouping as a tricarbonylmethane fragment, and the hydrogen chemical shift of the enol hydroxy group is $\sim 11$ p.p.m. (Wu et al., 2002). Most of the excellent inhibitors of p-hydroxyphenylpyruvate dioxygenase also possess similar characteristics, which are crucial for their two kinds of bioactivity (Zhu et al., 2004). Hitherto, we have synthesized a series of 3-(un)substituted benzoyl-1-alkyl-pyrrolidine-2,4-dione compounds, some of which have high herbicidal activity. The structure reported here, ( $1 b$ ), helps us to investigate the relationship between structure and herbicidal activity.


The analysis of crystals grown from a solution of 3-(2-methoxybenzoyl)-1-isopropylpyrrolidine-2,4-dione, (1a), showed that we had obtained crystals of the related tautomeric form, viz. 1-iso-propyl-3-( $\alpha$-hydroxy-2-methoxybenzylidene)-pyrrolidine-2,4-dione, $(1 b)$. The molecular structure of $(1 b)$ is shown in Fig. 1. Atom H2, involved in intramolecular hydrogen bonding between O 2 and O 4 , was assigned to O 2 rather than to O 4 , on the basis of the bond lengths. The $\mathrm{C} 11-$ O4 distance is 1.252 (3) $\AA$, which is longer than the normal carbonyl bond length ( $\mathrm{C} 9=\mathrm{O} 3$ ) of $1.210(5) \AA$. In contrast, the $\mathrm{C} 7-\mathrm{O} 2$ distance, 1.331 (3) $\AA$, is intermediate between the normal carbonyl $\mathrm{C}=\mathrm{O}$ bond and the $\mathrm{C}-\mathrm{O}$ single bond length (Allen et al., 1987). A similar situation has been reported for 3-(1-hydroxyethylidene)-1-phenylpyrrolidine-2,4-dione (Ellis \& Spek, 2001). The crystal structure of (1b) also involves two weak $\mathrm{C}-\mathrm{H} \cdots \mathrm{O}$ hydrogen-bonding interactions (Fig. 2 and Table 2).

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Figure 1
A view of the title compound, showing the atom-numbering scheme. Displacement ellipsoids are drawn at the $40 \%$ probability level. The intramolecular hydrogen bond is indicated by a dashed line.


Figure 2
A packing diagram, showing the intra- and intermolecular hydrogen bonds as dashed lines.

## Experimental

The title compound was obtained according to a reported procedure (Matsuo et al., 1980). Colorless single crystals were obtained by recrystallization of 1-isopropyl-3-( $\alpha$-hydroxy-2-methoxylbenzyl-idene)pyrrolidine-2,4-dione from petroleum ether and ethyl acetate.

## Crystal data

$\mathrm{C}_{15} \mathrm{H}_{17} \mathrm{NO}_{4}$
$M_{r}=275.30$
Orthorhombic, $P 2_{1} 2_{1} 2_{1}$
$a=6.981(2) \AA$
$b=13.720(4) \AA$
$c=14.863(5) \AA$
$V=1423.5(8) \AA^{3}$
$Z=4$
$D_{x}=1.285 \mathrm{Mg} \mathrm{m}^{-3}$

Mo $K \alpha$ radiation
Cell parameters from 939 reflections
$\theta=2.7-23.7^{\circ}$
$\mu=0.09 \mathrm{~mm}^{-1}$
$T=293$ (2) K
Prism, colorless
$0.22 \times 0.20 \times 0.16 \mathrm{~mm}$

## Data collection

| Bruker SMART CCD area-detector | 1701 independent reflections |
| :--- | :--- |
| $\quad$ diffractometer | 1300 reflections with $I>2 \sigma(I)$ |
| $\varphi$ and $\omega$ scans | $R_{\text {int }}=0.031$ |
| Absorption correction: multi-scan | $\theta_{\max }=26.4^{\circ}$ |
| $\quad(S A D A B S ;$ Sheldrick, 1996 $)$ | $h=-8 \rightarrow 8$ |
| $T_{\min }=0.965, T_{\max }=0.985$ | $k=-17 \rightarrow 14$ |
| 8260 measured reflections | $l=-18 \rightarrow 16$ |

## Refinement

Refinement on $F^{2}$
$R\left[F^{2}>2 \sigma\left(F^{2}\right)\right]=0.038$

$$
\begin{aligned}
& w=1 /[ \sigma^{2}\left(F_{o}^{2}\right)+(0.0464 P)^{2} \\
&+0.0951 P] \\
& \text { where } P=\left(F_{o}^{2}+2 F_{c}^{2}\right) / 3 \\
&(\Delta / \sigma)_{\max }<0.001 \\
& \Delta \rho_{\max }=0.11 \mathrm{e} \AA^{-3} \\
& \Delta \rho_{\min }=-0.14 \mathrm{e} \AA^{-3}
\end{aligned}
$$

$w R\left(F^{2}\right)=0.091$
$S=1.07$
1701 reflections
185 parameters
H -atom parameters constrained

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

| O1-C1 | 1.356 (3) | N1-C12 | 1.460 (3) |
| :---: | :---: | :---: | :---: |
| O1-C15 | 1.429 (3) | C6-C7 | 1.472 (3) |
| O2-C7 | 1.331 (3) | C7-C8 | 1.363 (3) |
| O3-C9 | 1.210 (3) | C8-C9 | 1.453 (3) |
| O4-C11 | 1.252 (3) | C8-C11 | 1.454 (3) |
| N1-C11 | 1.331 (3) | C9-C10 | 1.522 (4) |
| N1-C10 | 1.455 (3) |  |  |
| C1-O1-C15 | 118.1 (2) | O3-C9-C8 | 130.9 (2) |
| C11-N1-C10 | 111.63 (18) | $\mathrm{O} 3-\mathrm{C} 9-\mathrm{C} 10$ | 123.4 (2) |
| C11-N1-C12 | 124.8 (2) | C8-C9-C10 | 105.7 (2) |
| C10-N1-C12 | 123.5 (2) | $\mathrm{N} 1-\mathrm{C} 10-\mathrm{C} 9$ | 104.78 (19) |
| $\mathrm{O} 2-\mathrm{C} 7-\mathrm{C} 8$ | 119.5 (2) | O4-C11-N1 | 125.3 (2) |
| O2-C7-C6 | 113.0 (2) | $\mathrm{O} 4-\mathrm{C} 11-\mathrm{C} 8$ | 124.4 (2) |
| C8-C7-C6 | 127.5 (2) | N1-C11-C8 | 110.28 (18) |
| C7-C8-C9 | 131.4 (2) | N1-C12-C13 | 111.2 (2) |
| C7-C8-C11 | 120.8 (2) | N1-C12-C14 | 111.5 (2) |
| C9-C8-C11 | 107.6 (2) | C13-C12-C14 | 111.1 (3) |
| C5-C6-C7-O2 | -50.5 (3) | $\mathrm{O} 2-\mathrm{C} 7-\mathrm{C} 8-\mathrm{C} 11$ | -1.9 (3) |
| C1-C6-C7-O2 | 127.5 (2) | $\mathrm{C} 11-\mathrm{C} 8-\mathrm{C} 9-\mathrm{O} 3$ | 177.5 (3) |
| C5-C6-C7-C8 | 127.4 (3) | C7-C8-C9-C10 | -175.7 (2) |
| C1-C6-C7-C8 | -54.6 (3) | C12-N1-C10-C9 | 178.3 (2) |
| $\mathrm{O} 2-\mathrm{C} 7-\mathrm{C} 8-\mathrm{C} 9$ | 172.5 (3) | $\mathrm{O} 3-\mathrm{C} 9-\mathrm{C} 10-\mathrm{N} 1$ | -178.5 (2) |
| C6-C7-C8-C9 | -5.3 (4) |  |  |

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

| $D-\mathrm{H} \cdots A$ | $D-\mathrm{H}$ | $\mathrm{H} \cdots A$ | $D \cdots A$ | $D-\mathrm{H} \cdots A$ |
| :--- | :--- | :--- | :--- | :--- |
| $\mathrm{O} 2-\mathrm{H} 2 \cdots \mathrm{O} 4$ | 0.82 | 1.81 | $2.555(3)$ | 150 |
| $\mathrm{C} 15-\mathrm{H} 15 C \cdots \mathrm{O}^{\mathrm{i}}$ | 0.96 | 2.50 | $3.192(3)$ | 129 |
| $\mathrm{C}^{\mathrm{i}} 10-\mathrm{H} 10 A \cdots \mathrm{O}^{2}$ | 0.97 | 2.56 | $3.271(3)$ | 130 |

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

All H atoms were placed in calculated positions, with $\mathrm{C}-\mathrm{H}=0.93-$ $0.98 \AA$ and $\mathrm{O}-\mathrm{H}=0.82 \AA$, and included in the final cycles of refinement using a riding model, with $U_{\text {iso }}(\mathrm{H})=1.2 U_{\text {eq }}(\mathrm{C})$ or $1.5 U_{\text {eq }}(\mathrm{O})$. In the absence of significant anomalous dispersion effects, Friedel pairs were averaged, and the absolute configuration cannot be determined from the crystallographic experiment.

Data collection: SMART (Bruker, 1999); cell refinement: SMART; data reduction: SAINT (Bruker, 1999); program(s) used to solve structure: SHELXS 97 (Sheldrick, 1997); 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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## References

Allen, F. H., Kennard, O., Watson, D. G., Brammer, L., Orpen, A. G. \& Taylor, R. (1987). J. Chem. Soc. Perkin Trans. 2, pp. S1-19.

Bann, J. L. van der, Barnick, J. W. F. K. \& Bickelhaupt, F. (1978). Tetrahedron, 34, 223-231.
Bruker (1999). SMART, SAINT and SHELXTL. Bruker AXS Inc., Madison, Wisconsin, USA.
Ellis, D. D. \& Spek, A. L. (2001). Acta Cryst. C57, 433-434.
Holzapfel, C. W., Hutchison, R. D. \& Wilkins, D. C. (1970). Tetrahedron, 26, 5239-5246.
Matsuo, K., Kitaguchi, I., Takata, Y. \& Tanaka, K. (1980). Chem. Pharm. Bull. 28, 2494-2502.

Rinehart, K. L., Beck, J. R., Borders, D. B., Kinstle, T. H. \& Krauss, D. (1963). J. Am. Chem. Soc. 85, 4038-4039.

Rinehart, K. L., Mackellar, F. A., Grostic, M. F., Olson, E. C., Wnuk, R. J. \& Branfman, A. R. (1971). J. Am. Chem. Soc. 93, 4943-4945.
Rooyen, P. H. van (1992). Acta Cryst. C48, 551-552.
Sheldrick, G. M. (1996). SADABS. University of Göttingen, Germany.
Sheldrick, G. M. (1997). SHELSXS97 and SHELXL97. University of Göttingen, Germany.
Stickings, C. E. (1959). Biochem. J. 72, 332-334.
Wu, C.-S., Huang, J.-L., Sun, Y.-S. \& Yang, D.-Y. (2002). J. Med. Chem. 45, 2222-2228.
Zhu, Y.-Q., Hu, F.-Z. \& Yang, H.-Z. (2004). Huaxue Tongbao. In the press. (In Chinese.)

