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#### **Key indicators**

Single-crystal X-ray study T = 293 K Mean  $\sigma$ (C–C) = 0.002 Å R factor = 0.041 wR factor = 0.119 Data-to-parameter ratio = 12.8

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

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# Methyl 3-benzoyl-8-hydroxy-5-methoxyindolizine-1-carboxylate

In the title compound,  $C_{18}H_{15}NO_5$ , the methoxycarbonyl substituent, disregarding H atoms, is almost coplanar with the indolizine ring system in each of the two crystallographically independent molecules in the asymmetric unit. In the crystal structure, symmetry-related molecules form molecular dimers, held together by two  $C-H\cdots O$  hydrogen bonds, and are interconnected into a three-dimensional network by another  $C-H\cdots O$  hydrogen bond.

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#### Comment

Photoinduced oxygenation reactions of indolizine derivatives have been intensively investigated (Tian *et al.*, 2001). In continuation of this work, we prepared the title compound, (I), by photoinduced oxygenation of 1-acetyl-3-benzoylindolizine. As part of this work, we report here the crystal structure of (I).



The asymmetric unit of (I) consists of two crystallographically independent molecules, A and B. These are related by a local twofold axis, coupled with a shift of 2.15 Å parallel to this axis. The bond lengths and angles in the two molecules (Fig. 1) are in good agreement with each other, and the values are within normal ranges (Allen *et al.*, 1987).

In both molecules, A and B, the methoxycarbonyl groups (O3/O4/C9/C10) are almost coplanar with the indolizine ring system. The plane of the methoxycarbonyl group is twisted about the C8–C9 bond by an angle of 5.8 (1)° in molecule A and by an angle of 5.1 (1)° in molecule B. Coplanarity is maintained by an intramolecular O1–H1A···O3 hydrogen bond, linking the methoxycarbonyl and hydroxyl group (Fig. 1). The methoxy group (O2/C11), attached at atom C3, is slightly twisted out of the plane of the indolizine system, with a similar orientation in both molecules. The C4–C3–O2–C11 torsion angles are 24.3 (2) and 18.7 (3)° in molecules A and B, respectively, indicating that the methoxy group tends to deviate from coplanarity with the attached aromatic ring, as commonly observed in anisoles (Domiano *et al.*, 1979).

The relative configuration of the benzoyl substituent (O5/C12-C18) with respect to the indolizine is influenced by the

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The asymmetric unit of the title compound, showing 50% probability displacement ellipsoids and the atom-numbering scheme.



Figure 2

Packing diagram of the title compound, showing the formation of dimers.

 $sp^2$  hybridization of atom C12. In molecule A, the C2-C12-C13 angle is 121.3  $(1)^{\circ}$ , while the dihedral angle between the aromatic rings of the benzoyl group and the indolizine system is 64.6 (1)°. In molecule B, the corresponding values are 119.9 (1) and 61.8 (1)°, respectively. In the benzoyl substituent, atom O5 is out of the plane of the aromatic ring, such that the O5/C2/C12/C13 plane is twisted from the plane of the aromatic ring by an angle of 34.5  $(1)^{\circ}$  in molecule A and by an angle of 26.4  $(1)^{\circ}$  in molecule B.

In the crystal structure, two symmetry-related A molecules and two symmetry-related B molecules form molecular dimers (Fig. 2), by intermolecular C18-H18···O3 hydrogen bonds (Table 2). The dimers are interconnected into a threedimensional network by  $C11A - H11A \cdots O1B^{1}$  hydrogen bonds [symmetry code: (i) 2 - x, 1 - y, -z].

#### **Experimental**

The title compound was isolated from the reaction mixture of the photooxygenation of 1-acetyl-3-benzoylindolizine in methanol by column chromatography on silica gel. Single crystals, suitable for X- ray crystallographic measurements, were prepared by slow evaporation of the solvent from a petroleum ether-acetone solution (1:1 v/v).

5445 reflections with  $I > 2\sigma(I)$ 

 $w = 1/[\sigma^2(F_o^2) + (0.0611P)^2]$ 

where  $P = (F_o^2 + 2F_c^2)/3$ 

 $R_{\rm int} = 0.012$ 

 $\theta_{\rm max} = 27.0^{\circ}$ 

 $h = -11 \rightarrow 11$  $k = -14 \rightarrow 11$ 

 $l = -20 \rightarrow 11$ 

+ 0.333P]

 $(\Delta/\sigma)_{\rm max} < 0.001$  $\Delta \rho_{\rm max} = 0.19 \ {\rm e} \ {\rm \AA}^{-3}$ 

 $\Delta \rho_{\rm min} = -0.21 \text{ e } \text{\AA}^{-3}$ 

#### Crystal data

C <sub>18</sub> H <sub>15</sub> NO <sub>5</sub>	Z = 4
$M_r = 325.31$	$D_x = 1.397 \text{ Mg m}^{-3}$
Triclinic, P1	Mo $K\alpha$ radiation
$a = 9.0670 (4) \text{ Å}_{1}$	Cell parameters from 5941
b = 11.3774(5) Å	reflections
c = 15.9752 (8) Å	$\theta = 2.3 - 28.3^{\circ}$
$\alpha = 75.294 \ (1)^{\circ}$	$\mu = 0.10 \text{ mm}^{-1}$
$\beta = 76.127 \ (1)^{\circ}$	T = 293 (2)  K
$\gamma = 87.458 \ (1)^{\circ}$	Block, yellow
$V = 1547.26 (12) \text{ Å}^3$	$0.50 \times 0.50 \times 0.42 \text{ mm}$

#### Data collection

Siemens SMART CCD areadetector diffractometer (a) scans Absorption correction: none 9032 measured reflections 6518 independent reflections

#### Refinement

Refinement on  $F^2$  $R[F^2 > 2\sigma(F^2)] = 0.041$  $wR(F^2) = 0.120$ S = 1.036518 reflections 509 parameters H atoms treated by a mixture of independent and constrained refinement

#### Table 1

Selected geometric parameters (Å).

-			
O1A-C6A	1.3594 (16)	O1B-C6B	1.3536 (19)
N1A - C3A	1.3816 (16)	N1B-C3B	1.3825 (18)
N1A-C7A	1.3983 (14)	N1B-C7B	1.3951 (17)
N1A - C2A	1.4071 (16)	N1B-C2B	1.4021 (17)

Table 2	
TT	h

Hydrogen-bonding geometry (Å, °).

$D - H \cdots A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdots A$
$O1A - H1AA \cdots O3A$ $O1B - H1BA \cdots O3B$	0.93 (2) 0.95 (2)	1.71 (2) 1.68 (2)	2.629 (2) 2.614 (2)	175 (2) 165 (2)
$C11A - H11A \cdots O1B^{i}$	0.96	2.40	3.167 (2)	137
$C18A - H18A \cdots O3A^{ii}$	0.94 (2)	2.50 (2)	3.423 (2)	166 (2)
$C18B - H18B \cdots O3B^{iii}$	0.95 (2)	2.56 (2)	3.493 (2)	167 (2)

Symmetry codes: (i) 2 - x, 1 - y, -z; (ii) 1 - x, -y, 1 - z; (iii) 1 - x, 1 - y, -z.

The H atoms were located in difference Fourier maps and were refined isotropically, except for those of methyl groups C10 and C11, which were geometrically fixed and treated as riding atoms, with C-H distances of 0.96 Å and  $U_{iso}(H)$  vales equal to 1.5  $U_{eq}(C)$ . Due to a large fraction of weak data at higher angles, the  $2\theta$  maximum was limited to 54°.

Data collection: SMART (Siemens, 1996); cell refinement: SAINT (Siemens, 1996); data reduction: SAINT and SADABS (Sheldrick, 1996); program(s) used to solve structure: SHELXTL (Sheldrick, 1997); program(s) used to refine structure: SHELXTL; molecular graphics: SHELXTL; software used to prepare material for publication: SHELXTL, PARST (Nardelli, 1995) and PLATON (Spek, 1990).

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