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Key indicators

Single-crystal X-ray study T = 293 K Mean σ (C–C) = 0.004 Å R factor = 0.057 wR factor = 0.156 Data-to-parameter ratio = 15.2

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Dimethyl 7-(*N*,*N*-dimethylamino)-3-(4-methylbenzoyl)indolizine-1,2-dicarboxylate

In the title compound, $C_{22}H_{22}N_2O_5$, there are two molecules in the asymmetric unit. In each of them, the carboxylate groups are oriented perpendicular to each other and one of the carboxylate groups is almost coplanar with the indolizine moiety. In the solid state, weak intermolecular $C-H\cdots O$ contacts are observed. Received 16 February 2004 Accepted 2 March 2004 Online 13 March 2004

Comment

Chemists are attracted by indolizines and their derivatives because of their importance as pharmaceutical drugs, such as potential central nervous system depressants, cardiovascular agents, calcium entry blockers, spectral sensitizers and novel dyes (Hema *et al.*, 2003, and references therein). Indolizine derivatives such as 1-carboxymethyl-3-(4-chlorobenzoyl)-7methoxy-2-methylindolizine, 3-acetyl(benzoyl)-1-carboxyethyl)indolizine and 3-carboxymethyl-1-(4-chlorobenzoyl)-7methoxy-2-methylindolizine exhibit *anti*-inflammatory activity (Casagrande *et al.*, 1971). 3–Benzoylindolizine-1-acetic acid exhibits an auxin-like activity (Carbellini *et al.*, 1968). In view of the these important attributes, the structure determination of the title compound, (I), was performed.



The asymmetric unit of (I) contains two crystallographically independent molecules (A and B). A view of the two independent molecules, including the atomic numbering scheme, is shown in Fig. 1. The overall weighted r.m.s. fit of molecules A and B is 0.095 Å. The corresponding bond lengths and angles in the independent molecules agree with each other and are comparable to those in related structures (Hema *et al.*, 2003, 2004). The planes of the 1- and 2-carboxylate groups are oriented at angles of 4.52 (6) [3.15 (10)° for molecule B] and 72.05 (7)° [74.67 (14)° for molecule B], respectively, with respect to the plane of the indolizine moiety. The corresponding dihedral angles in a related structure (Hema *et al.*, 2004) are 5.97 (9) and 72.05 (7)°, respectively. The carboxylate groups are oriented perpendicular to each other. The dihedral angle between the planes of the indolizine moiety and the



The two independent molecules of (I), with the atom-labelling scheme. Displacement ellipsoids are drawn at the 30% probability level. H atoms are represented by circles of arbitrary radii.

benzoyl ring is 59.13 (6)° [58.31 (6)° for molecule B], while the corresponding angle in a related structure (Hema et al., 2004) is 58.13 (5)°.

In the crystal structure, atom C5B of molecule B acts as a donor for a weak intermolecular C-H···O interaction with carbonyl atom O16B of a symmetry-related molecule at (1 - x, -y, -z), leading to an $R_2^2(12)$ motif (Bernstein *et al.*, 1995). Atom C15B of molecule B is involved in a weak intermolecular C-H···O interaction with atom O10B of a symmetry-related molecule at (2 - x, 1 - y, 1 - z) and has a graph-set motif of $R_2^2(16)$ (Table 1). Atom C25B acts as a donor for a weak intermolecular C-H···O interaction with carbonyl atom O16A of molecule A of an adjacent molecule.

Experimental

A mixture of 4-dimethylaminopyridinium-1-(4-methyl)phenacylide (1.4 mmol), dimethyl acetylenedicarboxylate (1.6 mmol) and potassium carbonate (1.6 mmol) in dimethylformamide (30 ml) was kept at room temperature overnight. The insoluble materials were removed by filtration, and the filtrate was extracted with an ethyl acetate/dilute HCl mixture. The organic layer was evaporated and chromatographed to give (I), which was recrystallized from ethyl acetate (yield 47%, m.p. 443-445 K).

Z = 4

 $D_x = 1.308 \text{ Mg m}^{-3}$

Cell parameters from 3068

Mo $K\alpha$ radiation

reflections

 $\mu = 0.09 \text{ mm}^{-1}$

T = 293 (2) K

Prism, yellow $0.28 \times 0.20 \times 0.15 \text{ mm}$

 $\theta=2.4{-}24.5^\circ$

Crystal data

$C_{22}H_{22}N_2O_5$
$M_r = 394.42$
Triclinic, P1
a = 8.050 (5) Å
b = 16.942 (11) Å
c = 17.094 (11) Å
$\alpha = 118.555 \ (10)^{\circ}$
$\beta = 95.033 \ (12)^{\circ}$
$\gamma = 97.082 \ (11)^{\circ}$
$V = 2003 (2) \text{ Å}^3$

 ω scans Absorption correction: none

Data collection

Siemens SMART CCD area-

detector diffractometer

 $h=-10\to9$ $k = -22 \rightarrow 17$ 12 347 measured reflections $l = -22 \rightarrow 22$ 8081 independent reflections Refinement Refinement on F^2 $w = 1/[\sigma^2(F_o^2) + (0.0707P)^2]$ $R[F^2 > 2\sigma(F^2)] = 0.057$ + 0.2476P] $wR(F^2) = 0.156$ where $P = (F_0^2 + 2F_c^2)/3$ S=1.03 $(\Delta/\sigma)_{\rm max} < 0.001$ $\Delta \rho_{\rm max} = 0.17 \ {\rm e} \ {\rm \AA}^{-3}$ 8081 reflections $\Delta \rho_{\rm min} = -0.21 \text{ e} \text{ Å}^{-3}$ 533 parameters

5203 reflections with $I > 2\sigma(I)$

 $R_{\rm int} = 0.018$

 $\theta_{\rm max} = 28.1^\circ$

Table 1

 $C-H \cdots O$ interactions (Å, °).

H-atom parameters constrained

$D - H \cdot \cdot \cdot A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - H \cdots A$
$C5B-H52\cdots O16B^{i}$ $C15B-H155\cdots O10B^{ii}$	0.93 0.96	2.45 2.52	3.154 (3) 3.417 (4)	132 156
$C25B - H256 \cdots O16A^{iii}$	0.96	2.60	3.531 (4)	164

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

The methyl H atoms were constrained to an ideal geometry (C-H = 0.96 Å), with U_{iso} (H) values of $1.5U_{iso}$ (C), but were allowed to rotate freely about the C-C bond. All remaining H atoms were placed in idealized positions (C–H = 0.93 Å) and constrained to ride on their parent atoms, with $U_{iso}(H)$ values of $1.2U_{iso}(C)$.

Data collection: SMART (Siemens, 1996); cell refinement: SAINT (Siemens, 1996); data reduction: SAINT; program(s) used to solve structure: SIR92 (Altomare et al., 1994); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: ORTEP-3 (Farrugia, 1997); software used to prepare material for publication: SHELXL97 and PLATON (Spek, 2003).

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