

(Z)-3-Acetylidene-2,3-dihydro-1H-isoindolin-1-one

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Key indicators

Single-crystal X-ray study

$T = 293\text{ K}$

Mean $\sigma(\text{C}-\text{C}) = 0.002\text{ \AA}$

R factor = 0.043

wR factor = 0.121

Data-to-parameter ratio = 19.1

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

The asymmetric unit of the title compound, $\text{C}_{11}\text{H}_9\text{NO}_2$, contains two molecules which are related by a pseudo-inversion centre and form an $\text{N}-\text{H}\cdots\text{O}$ hydrogen-bonded dimer. The least-squares planes through the non-H atoms of the molecules illustrate the overall molecular planarity. The molecules assume *Z* configurations about the exocyclic $\text{C}=\text{C}$ bond. In the crystal, intermolecular hydrogen bonds link the molecules into infinite chains.

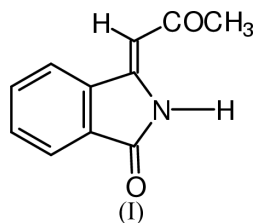
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Comment

The isoindolinone skeleton is present in a number of natural products (Tamaoki *et al.*, 1986; Valencia *et al.*, 1985). Many isoindolinone derivatives exhibit biological activity as potential antiviral, antileukemic, anti-inflammatory, antipsychotic and antiulcer agents (Zhuang *et al.*, 1998; Taylor *et al.*, 1997; De Clercq, 1995; Pendrak *et al.*, 1994). As part of our ongoing studies on the synthesis and characterization of new heterocyclic systems containing an isoindolinone moiety (Mukherjee *et al.*, 2000; Kundu, Khan & Mukhopadhyay, 1999), the structure determination of (*Z*)-3-acetylidene-2,3-dihydro-1H-isoindolin-1-one, (**I**), was undertaken.



The two approximately parallel molecules (*A* and *B*) in the asymmetric unit related by a pseudo-inversion centre at (0.50, 0.38, 0.25), can be described as an $\text{N}-\text{H}\cdots\text{O}$ bonded dimer (Fig. 1). The dihedral angle between the nearly planar molecules *A* and *B* is $2.45(2)^\circ$. The *Z* configuration of the molecules is established by the $\text{N}1-\text{C}8-\text{C}9-\text{C}10$ torsion angles of $-4.6(2)^\circ$ in *A* and $4.3(2)^\circ$ in *B*. The angles between the five- and six-membered rings of the isoindolinone system in *A* and *B* are $1.92(6)^\circ$ and $1.54(6)^\circ$, respectively. The bond distances and angles in (**I**) are similar to those reported for substituted isoindoline structures (Kundu, Khan, Guha & Mukherjee, 1999; Khan *et al.*, 1998; McNab *et al.*, 1997). In the solid state, weak intermolecular $\text{C}9-\text{H}9\cdots\text{O}1$ hydrogen bonds link the dimers to form infinite chains which are cross-linked in the *c* direction by weak intermolecular $\text{C}4\text{B}-\text{H}4\text{B}\cdots\text{O}2\text{A}$ hydrogen bonds (Table 2), to form a network.

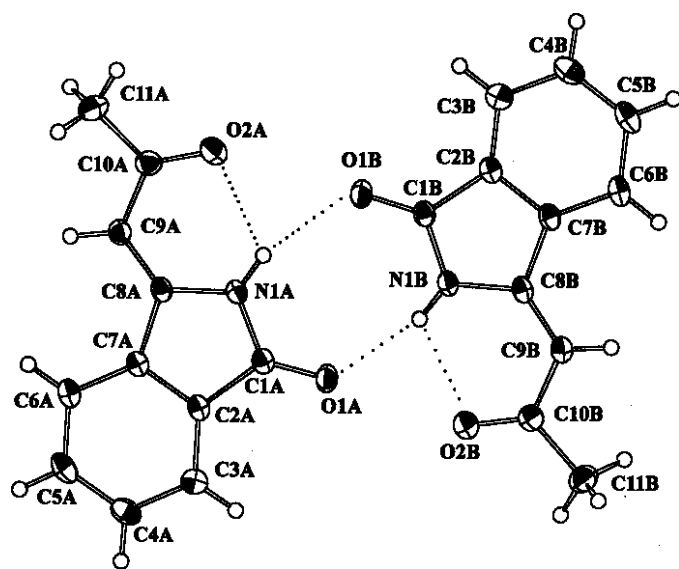


Figure 1
ZORTEP view (50% probability level) of the two hydrogen-bonded molecules of (I).

Experimental

2-[(Trimethylsilyl)ethynyl]benzamide (TEB) was prepared by stirring a mixture of *o*-iodobenzamide (1 mmol), (trimethylsilyl)acetylene (2 mmol), $(\text{PPh}_3)_2\text{PdCl}_2$ (3.5 mol%), CuI (8 mol%) and triethylamine (4 mmol) in dimethylformamide at room temperature under a nitrogen atmosphere for 24 h. The reaction of TEB (1 mmol) with CH_3COCl (acetyl chloride, 1.2 mmol) in the presence of anhydrous AlCl_3 (4 mmol) and tetrachloroethane (10 ml) at 273 K yielded the title compound, (I) (m.p. 381 K). Single crystals suitable for X-ray analysis were obtained by slow crystallization from a dilute solution of (I) in ethanol. Elemental analysis, calculated for $\text{C}_{11}\text{H}_9\text{NO}_2$: C 70.58, H 4.81, N 7.48%; found: C 70.53, H 4.89, N 7.24%.

Crystal data

$\text{C}_{11}\text{H}_9\text{NO}_2$
 $M_r = 187.19$
 Monoclinic, $P2_1/n$
 $a = 6.8673$ (2) Å
 $b = 12.8243$ (3) Å
 $c = 20.6130$ (5) Å
 $\beta = 94.520$ (1)°
 $V = 1809.71$ (8) Å³
 $Z = 8$

$D_x = 1.374$ Mg m⁻³
 Mo $K\alpha$ radiation
 Cell parameters from 922 reflections
 $\theta = 5.4\text{--}31.7^\circ$
 $\mu = 0.10$ mm⁻¹
 $T = 293$ (2) K
 Hexagonal block, colourless
 $0.50 \times 0.45 \times 0.30$ mm

Data collection

Bruker SMART-CCD diffractometer
 ω scans
 14 180 measured reflections
 4883 independent reflections
 4172 reflections with $I > 2\sigma(I)$

Refinement

Refinement on F^2
 $R[F^2 > 2\sigma(F^2)] = 0.043$
 $wR(F^2) = 0.121$
 $S = 1.05$
 4883 reflections
 255 parameters
 H-atom parameters constrained

$R_{\text{int}} = 0.030$
 $\theta_{\text{max}} = 29.5^\circ$
 $h = -9 \rightarrow 9$
 $k = -17 \rightarrow 17$
 $l = -19 \rightarrow 27$

$w = 1/[\sigma^2(F_o^2) + (0.0603P)^2 + 0.6557P]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\text{max}} < 0.001$
 $\Delta\rho_{\text{max}} = 0.36$ e Å⁻³
 $\Delta\rho_{\text{min}} = -0.26$ e Å⁻³

Table 1

Selected geometric parameters (Å, °).

N1A—C1A	1.3842 (15)	N1B—C1B	1.3863 (15)
N1A—C8A	1.3931 (14)	N1B—C8B	1.3923 (13)
C7A—C8A	1.4768 (15)	C7B—C8B	1.4757 (15)
C8A—C9A	1.3513 (16)	C8B—C9B	1.3512 (16)
C9A—C10A	1.4682 (16)	C9B—C10B	1.4682 (16)
C1A—N1A—C8A	112.57 (9)	C1B—N1B—C8B	112.68 (9)
C9A—C8A—N1A	126.67 (10)	C9B—C8B—N1B	126.87 (11)
C9A—C8A—C7A	127.24 (10)	C9B—C8B—C7B	127.19 (10)
N1A—C8A—C7A	105.98 (9)	N1B—C8B—C7B	105.90 (9)
C8A—C9A—C10A	123.37 (11)	C8B—C9B—C10B	123.66 (11)
N1A—C8A—C9A—C10A	-4.56 (18)	N1B—C8B—C9B—C10B	4.26 (18)
C7A—C8A—C9A—C10A	171.14 (10)	C7B—C8B—C9B—C10B	-172.90 (10)

Table 2

Hydrogen-bonding geometry (Å, °).

$D\cdots H\cdots A$	$D\cdots H$	$H\cdots A$	$D\cdots A$	$D\cdots H\cdots A$
N1A—H1A ⁱ ···O2A	0.86	2.25	2.776 (1)	120
N1A—H1A ⁱ ···O1B	0.86	2.38	3.065 (1)	137
N1B—H1B ⁱⁱ ···O1A	0.86	2.30	2.977 (1)	135
N1B—H1B ⁱⁱ ···O2B	0.86	2.26	2.787 (1)	120
C9A—H9A ⁱ ···O1A ⁱ	0.93	2.49	3.208 (1)	135
C4B—H4B ⁱⁱⁱ ···O2A ⁱⁱⁱ	0.93	2.58	3.488 (2)	167
C9B—H9B ⁱⁱⁱ ···O1B ⁱⁱⁱ	0.93	2.57	3.283 (1)	134

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

The H atoms were refined using a riding model and their isotropic displacement parameters were set to 1.2 times (1.5 times for CH_3 groups) the equivalent displacement parameters of their parent atoms.

Data collection: SMART (Siemens, 1996); cell refinement: SMART; data reduction: SAINT (Siemens, 1996); program(s) used to solve structure: MULTAN88 (Debaerdemaeker *et al.*, 1988); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: ZORTEP (Zsolnai, 1995); software used to prepare material for publication: SHELXL97).

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