organic papers

Acta Crystallographica Section E Structure Reports Online

ISSN 1600-5368

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Key indicators

Single-crystal X-ray study T = 292 KMean $\sigma(\text{C}-\text{C}) = 0.005 \text{ Å}$ Disorder in main residue R factor = 0.048 wR factor = 0.129 Data-to-parameter ratio = 12.9

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

2-(5-Benzyloxy-2,4-dichlorophenyl)-4,5,6,7-tetrahydroisoindole-1,3-dione

The crystal structure of the title compound, $C_{21}H_{17}Cl_2NO_3$, shows that there are no intra- or intermolecular π - π stacking interactions. The structure is stabilized by C-H···O and C-H···Cl hydrogen bonds, which form a nine-membered ring.

Comment

It has been found that some cyclic imides show strong herbicidal activity at small doses against a wide variety of weeds including Gramineae weeds, Cyperaceae weeds and broadleaved weeds in upland fields, as well as weeds in paddy fields, and these compounds do not produce any material phytotoxicity on agricultural crops such as corn, soybean, wheat, peanut and rice plants (Nagano *et al.*, 1982). We are interested in exploring and developing novel strategies for synthesizing cyclic imides. The title compound, (I), was synthesized directly from the hydroxy analog.



In the molecule (Fig. 1), the dihedral angle between the phenyl (C1–C6) and the benzene (C8–C13) rings is 101.6 (1)°. The dihedral angle between the phenyl ring and the fivemembered heterocyclic ring (C14–C16/C21/N1) is 37.9 (1)°. The dihedral angle between the benzene and heterocyclic rings is 85.6 (1)°. There are intermolecular C–H···O and C– H···Cl hydrogen bonds (Table 1), which form a ninemembered ring, as shown in Fig. 2.

Experimental

5-Benzyloxy-2,4-dichlorophenylamine (0.14 g) and 3,4,5,6-tetrahydrophthalic anhydride (0.28 g) were dissolved in acetic acid (10 ml) and refluxed for 4 h. The resultant mixture was allowed to cool to room temperature and poured into water, followed by extraction with diethyl ether. This extract was washed with water, dried over anhydrous sodium sulfate and subjected to filtration. The filtrate was concentrated under reduced pressure and the residue was purified by silica gel chromatography to obtain (I) (0.30 g). Single crystals suitable for X-ray diffraction were grown from an ethanol solution at 277 K.

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Accepted 27 June 2005

Online 6 July 2005



Figure 1

A view of the molecule of (I), showing the atom-labeling scheme. Displacement ellipsoids are drawn at the 50% probability level. H atoms have been omitted for clarity. Dashed lines denote the minor component of the disorder.

Crystal data

 $C_{21}H_{17}Cl_2NO_3$ $M_r = 402.26$ Monoclinic, $P2_1/c$ a = 12.1705 (17) Å b = 11.0375 (15) Å c = 15.078 (2) Å $\beta = 107.232$ (3)° V = 1934.6 (5) Å³ Z = 4

Data collection

Bruker SMART CCD area-detector2169 reflediffractometer $R_{int} = 0.02$ φ and ω scans $\theta_{max} = 25$ Absorption correction: noneh = -149530 measured reflectionsk = -103386 independent reflectionsl = -17

Refinement

Refinement on F^2 $R[F^2 > 2\sigma(F^2)] = 0.048$ $wR(F^2) = 0.129$ S = 1.053386 reflections 263 parameters H-atom parameters constrained 2169 reflections with $I > 2\sigma(I)$ $R_{\text{int}} = 0.038$ $\theta_{\text{max}} = 25.0^{\circ}$ $h = -14 \rightarrow 9$ $k = -10 \rightarrow 13$ $l = -17 \rightarrow 17$

 $D_r = 1.381 \text{ Mg m}^{-3}$

Cell parameters from 1743

Mo Ka radiation

reflections

 $\begin{array}{l} \theta = 2.3 {-} 20.4^{\circ} \\ \mu = 0.36 \ \mathrm{mm}^{-1} \end{array}$

T = 292 (2) K

Plate, colorless $0.30 \times 0.20 \times 0.08 \text{ mm}$

$$\begin{split} w &= 1/[\sigma^2(F_{\rm o}^{-2}) + (0.0548P)^2 \\ &+ 0.1448P] \\ \text{where } P &= (F_{\rm o}^2 + 2F_{\rm c}^{-2})/3 \\ (\Delta/\sigma)_{\rm max} &< 0.001 \\ \Delta\rho_{\rm max} &= 0.23 \text{ e} \text{ Å}^{-3} \\ \Delta\rho_{\rm min} &= -0.18 \text{ e} \text{ Å}^{-3} \end{split}$$

Table 1 Hydrogen-bond geometry (Å °)

Trydrogen-bond	geometry	(A,).

$D - H \cdots A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdot \cdot \cdot A$
$C20-H20A\cdots Cl1^{i}$	0.97	2.81	3.745 (4)	163
$C10-H10\cdots O2^{ii}$	0.93	2.51	3.393 (4)	159
Symmetry codes: (i) $-x$	$+1, y + \frac{1}{2}, -z$	$+\frac{1}{2}$; (ii) $-x + 1$	$, y - \frac{1}{2}, -z + \frac{1}{2}.$	

H atoms were placed at calculated positions and treated as riding atoms, with C–H = 0.93 or 0.97 Å, and $U_{\rm iso}(\rm H) = 1.2 U_{eq}(\rm C)$. Atoms C18 and C19 were described by two disorder components each, with separately refined displacement parameters. The site-occupancy factors of the disordered atoms refined to 0.571 (15) and 0.429 (15).

Data collection: *SMART* (Bruker, 1997); cell refinement: *SAINT* (Bruker, 1999); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine



Figure 2

Hydrogen bonding in the crystal structure of (I). Hydrogen bonds are shown as dashed lines. [Symmetry codes: (i) 1 - x, $\frac{1}{2} + y$, $\frac{1}{2} - z$; (ii) 1 - x, $-\frac{1}{2} + y$, $\frac{1}{2} - z$].

structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *SHELXTL* (Bruker, 2001); software used to prepare material for publication: *SHELXTL*.

The authors acknowledge financial support from the National Key Project for Basic Research (2002CCA00500), the National Natural Science Foundation of China (No. 20432010, 20476036 and 20172017), the Program for New Century Excellent Talents in Universities of China and the Program for Excellent Research Groups of Hubei Province (No. 2004ABC002).

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