Acta Crystallographica Section E

## Structure Reports

Online
ISSN 1600-5368

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## Key indicators

Single-crystal X-ray study
$T=294 \mathrm{~K}$
Mean $\sigma(\mathrm{C}-\mathrm{C})=0.003 \AA$
$R$ factor $=0.044$
$w R$ factor $=0.115$
Data-to-parameter ratio $=18.6$

For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e.
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# $N^{\prime}$-(2,4-Dichlorobenzylidene)benzohydrazide 

The asymmetric unit of the title compound, $\mathrm{C}_{14} \mathrm{H}_{10} \mathrm{Cl}_{2} \mathrm{~N}_{2} \mathrm{O}$, comprises two independent molecules, in which the dihedral angles between the terminal phenyl and substituted benzene rings are 21.28 (10) and $39.60(12)^{\circ} . \mathrm{N}-\mathrm{H} \cdots \mathrm{O}, \mathrm{C}-\mathrm{H} \cdots \mathrm{O}$ and $\mathrm{C}-\mathrm{H} \cdots \mathrm{Cl}$ hydrogen bonds stabilize the conformations of the molecules and also stabilize the structure in the solid state.

## Comment

Metal complexes based on Schiff bases have attracted much attention because they can be utilized as model compounds of active centres in various proteins and enzymes (Kahwa et al., 1986; Santos et al., 2001). We report the synthesis and structure of the title compound, (I), as part of an investigation of its physical and chemical properties (Yu et al., 2005; Deng et al., 2005; Jing et al., 2005a, 2005b).

(I)

The asymmetric unit comprises two independent molecules (Fig. 1 and Table 1). In molecule 1, the dihedral angle between the terminal phenyl (C2-C7) and benzene (C9-C14) rings is $21.28(10)^{\circ}$, while in molecule 2 the corresponding angle is 39.60 (12). In the crystal structure, $\mathrm{N}-\mathrm{H} \cdots \mathrm{O}, \mathrm{C}-\mathrm{H} \cdots \mathrm{O}$ and $\mathrm{C}-\mathrm{H} \cdots \mathrm{Cl}$ hydrogen bonds stabilize the conformations of the molecules and also stabilize the structure in the solid state (Table 2 and Fig. 2).


Figure 1
The structure of the asymmetric unit of (I), with displacement ellipsoids for non-H atoms drawn at the $30 \%$ probability level.

Received 27 October 2005 Accepted 21 November 2005 Online 26 November 2005


Figure 2
Packing view of (I), viewed down the $a$ axis, showing intermolecular hydrogen bonds (dashed lines).

## Experimental

An anhydrous ethanol solution of 2,4-dichlorobenzaldehyde (1.75 g, 10 mmol ) was added to an anhydrous ethanol solution of benzohydrazide ( $1.36 \mathrm{~g}, 10 \mathrm{mmol}$ ) and the mixture was stirred at 350 K for 5 h under nitrogen, whereupon a precipitate appeared. The product was isolated and recrystallized from ethanol, and then dried in vacuo to give pure (I) in $89 \%$ yield. Colorless single crystals of (I) suitable for X-ray analysis were obtained by slow evaporation of an ethanol solution.

## Crystal data

## $\mathrm{C}_{14} \mathrm{H}_{10} \mathrm{Cl}_{2} \mathrm{~N}_{2} \mathrm{O}$ <br> $M_{r}=293.14$ <br> Monoclinic, $P 2_{1} / n$ <br> $a=10.5668$ (9) $\AA$ <br> $b=17.8266$ (15) Å <br> $c=14.9814$ (13) $\AA$ <br> $\beta=104.734$ (1) ${ }^{\circ}$ [please check] <br> $V=2729.3$ (4) $\AA^{3}$ <br> $Z=8$

## $D_{x}=1.427 \mathrm{Mg} \mathrm{m}^{-3}$

Mo $K \alpha$ radiation
Cell parameters from 3851

## reflections

$\theta=2.3-24.7^{\circ}$
$\mu=0.47 \mathrm{~mm}^{-1}$
$T=294$ (2) K
Prism, colorless
$0.46 \times 0.34 \times 0.22 \mathrm{~mm}$

## Data collection

Bruker SMART CCD area-detector diffractometer
$\varphi$ and $\omega$ scans
Absorption correction: multi-scan (SADABS; Sheldrick, 1996) $T_{\text {min }}=0.822, T_{\text {max }}=0.906$
18247 measured reflections

## Refinement

Refinement on $F^{2}$

$$
\begin{aligned}
& w=1 /\left[\sigma^{2}\left(F_{\mathrm{o}}{ }^{2}\right)+(0.0438 P)^{2}\right. \\
& +0.8233 P] \\
& \text { where } P=\left(F_{\mathrm{o}}{ }^{2}+2 F_{\mathrm{c}}{ }^{2}\right) / 3 \\
& (\Delta / \sigma)_{\max }=0.001 \\
& \Delta \rho_{\text {max }}=0.46 \mathrm{e}^{-3}{ }^{-3} \\
& \Delta \rho_{\text {min }}=-0.49 \mathrm{e}^{-3} \\
& \text { Extinction correction: SHELXL97 } \\
& \text { Extinction coefficient: } 0.0027 \text { (4) }
\end{aligned}
$$

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

| C1-N1-N2 | $119.27(16)$ | C22-N4-N3 | $115.67(16)$ |
| :--- | :---: | :--- | :---: |
| C8-N2-N1 | $115.09(16)$ | O1-C1-N1 | $121.40(17)$ |
| C15-N3-N4 | $117.99(16)$ | O2-C15-N3 | $122.30(18)$ |
|  |  |  |  |
| C1-N1-N2-C8 | $-179.31(18)$ | $\mathrm{N} 1-\mathrm{N} 2-\mathrm{C} 8-\mathrm{C} 9$ | $176.43(16)$ |
| C15-N3-N4-C22 | $177.48(18)$ | $\mathrm{N} 4-\mathrm{N} 3-\mathrm{C} 15-\mathrm{O} 2$ | $8.9(3)$ |
| $\mathrm{N} 2-\mathrm{N} 1-\mathrm{C} 1-\mathrm{O} 1$ | $-8.3(3)$ | $\mathrm{N} 4-\mathrm{N} 3-\mathrm{C} 15-\mathrm{C} 16$ | $-169.43(17)$ |
| $\mathrm{N} 2-\mathrm{N} 1-\mathrm{C} 1-\mathrm{C} 2$ | $170.93(16)$ | $\mathrm{N} 3-\mathrm{N} 4-\mathrm{C} 22-\mathrm{C} 23$ | $-174.83(17)$ |

Table 2
Hydrogen-bond geometry ( $\AA^{\circ}{ }^{\circ}$ ).

| $D-\mathrm{H} \cdots A$ | $D-\mathrm{H}$ | $\mathrm{H} \cdots A$ | $D \cdots A$ | $D-\mathrm{H} \cdots A$ |
| :---: | :---: | :---: | :---: | :---: |
| $\mathrm{N} 3-\mathrm{H} 3 \cdots \mathrm{O} 1^{\text {i }}$ | 0.88 (2) | 2.20 (2) | 3.007 (2) | 153 (2) |
| $\mathrm{N} 1-\mathrm{H} 1 \cdots \mathrm{O} 2$ | 0.86 (2) | 2.16 (2) | 3.005 (2) | 169 (2) |
| $\mathrm{C} 7-\mathrm{H} 7 \cdots \mathrm{O} 2$ | 0.93 | 2.45 | 3.372 (3) | 172 |
| $\mathrm{C} 8-\mathrm{H} 8 \cdots \mathrm{O} 2$ | 0.93 | 2.52 | 3.332 (2) | 146 |
| $\mathrm{C} 11-\mathrm{H} 11 \cdots \mathrm{O} 1^{\text {ii }}$ | 0.93 | 2.59 | 3.522 (3) | 178 |
| $\mathrm{C} 13-\mathrm{H} 13 \cdots \mathrm{Cl} 2^{\text {iii }}$ | 0.93 | 2.74 | 3.605 (2) | 156 |
| $\mathrm{C} 17-\mathrm{H} 17 \cdots \mathrm{O}^{1}$ | 0.93 | 2.50 | 3.416 (3) | 170 |
| Symmetry codes: $\begin{equation*} x+\frac{1}{2},-y+\frac{1}{2}, z+\frac{1}{2} . \tag{iii} \end{equation*}$ | (i) $x-\frac{1}{2},-y+\frac{1}{2}, z-\frac{1}{2}$; <br> (ii) $x+\frac{1}{2},-y+\frac{1}{2}, z-\frac{1}{2}$; |  |  |  |

H atoms attached to N atoms were located in a difference Fourier map and refined freely. Other H atoms were included in calculated positions $(\mathrm{C}-\mathrm{H}=0.93-0.96 \AA)$ and refined using a riding model approximation, with $U_{\text {iso }}(\mathrm{H})=1.2 U_{\mathrm{eq}}(\mathrm{C})$ or $1.5 U_{\mathrm{eq}}$ (methyl C).

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