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Key indicators

Single-crystal X-ray study T = 298 K Mean σ (C–C) = 0.003 Å R factor = 0.035 wR factor = 0.079 Data-to-parameter ratio = 14.2

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

2'-[1-(3,5-Dichloro-2-hydroxyphenyl)methylidene]isonicotinohydrazide monohydrate

The asymmetric unit of the title compound, $C_{13}H_9Cl_2N_3O_{2}$ - H_2O , consists of a Schiff base molecule and a solvent water molecule. The Schiff base molecule is approximately planar and displays a *trans* configuration with respect to the C=N double bond. The crystal structure is stabilized by intermolecular N-H···O, O-H···O, C-H···N, and C-H···O hydrogen bonds, forming a network.

Comment

Schiff base compounds have been of great interest for a long time and play an important role in the development of coordination chemistry (Musie *et al.*, 2001; Bernardo *et al.*, 1996; Paul *et al.*, 2002). As an extension of our work on the structural characterization of such compounds (Yang, 2006*a*,*b*,*c*), the crystal structure of the title compound, (I), is reported here.



The asymmetric unit of (I) comprises a Schiff base molecule and a solvent water molecule (Fig. 1), in which all the bond lengths are within normal ranges (Allen *et al.*, 1987) and comparable with those of similar compounds (Qian *et al.*, 2006; Qiu, Xu *et al.*, 2006; Qiu, Fang, Liu & Zhu, 2006; Qiu, Fang, Yang *et al.*, 2006; Yang, 2006*d*,*e*). The C7=N1 bond length of 1.278 (3) Å confirms it as a double bond, with a *trans* configuration. The C8-N2 bond is also relatively short [1.358 (3) Å], suggesting some degree of delocalization in the acetohydrazide system. The dihedral angle between the benzene ring and the pyridine ring is 5.6 (2)°.

In the crystal structure, water molecules are linked to the Schiff base molecules through $O-H\cdots O$ hydrogen bonds (Table 1). The molecules are further linked through intermolecular $N-H\cdots O$, $C-H\cdots N$ and $C-H\cdots O$ hydrogen bonds, forming a network (Fig. 2).

Experimental

3,5-Dichlorosalicylaldehyde (0.1 mmol, 19.2 mg) and pyridine-4carboxylic acid hydrazide (0.1 mmol, 13.7 mg) were dissolved in MeOH–H₂O (10 ml, 10:1 ν/ν). The mixture was stirred at room temperature to give a clear yellow solution. Crystals of the title Received 31 August 2006 Accepted 4 September 2006

04414 Yang and Guo $\cdot C_{13}H_9Cl_2N_3O_2H_2O$

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Figure 1

The molecular structure of (I), showing the atom-numbering scheme. Displacement ellipsoids are drawn at the 30% probability level. An intramolecular hydrogen bond is shown as a dashed line.

compound were formed by gradual evaporation of the solvent over a period of about one week at room temperature. Analysis, found: C 47.40, H 3.49, N 12.73%; calculated for $C_{13}H_{11}Cl_2N_3O_3$: C 47.58, H 3.38, N 12.81%.

Z = 2

 $D_x = 1.558 \text{ Mg m}^{-3}$

 $0.27 \times 0.20 \times 0.08 \text{ mm}$

5674 measured reflections

2831 independent reflections 2544 reflections with $I > 2\sigma(I)$

Mo $K\alpha$ radiation

 $\mu = 0.48 \text{ mm}^{-1}$

T = 298 (2) K Plate, yellow

 $R_{\rm int} = 0.023$

 $\theta_{\rm max} = 26.5^\circ$

Crystal data

$C_{13}H_9Cl_2N_3O_2{\cdot}H_2O$
$M_r = 328.15$
Monoclinic, P21
a = 3.887 (1) Å
<i>b</i> = 13.638 (2) Å
c = 13.231 (2) Å
$\beta = 94.546 \ (2)^{\circ}$
$V = 699.2 (2) \text{ Å}^3$

Data collection

Bruker SMART CCD area-detector diffractometer ω scans Absorption correction: multi-scan (*SADABS*; Sheldrick, 1996) $T_{\min} = 0.882, T_{\max} = 0.963$

Refinement

Refinement on F^2 $w = 1/[\sigma^2(F_0^2) + (0.0386P)^2]$ $R[F^2 > 2\sigma(F^2)] = 0.035$ wR(F²) = 0.079 where $P = (F_0^2 + 2F_c^2)/3$ $(\Delta/\sigma)_{\rm max} < 0.001$ $\Delta \rho_{\rm max} = 0.20 \text{ e } \text{\AA}^{-3}$ S = 1.02 $\Delta \rho_{\rm min} = -0.18 \text{ e } \text{\AA}^{-3}$ 2831 reflections 200 parameters Absolute structure: Flack (1983), H atoms treated by a mixture of 1334 Friedel pairs independent and constrained Flack parameter: 0.02 (6) refinement

Table 1

Hydrogen-bond geometry (Å, °).

$D - H \cdots A$	D-H	$H \cdots A$	$D \cdots A$	$D - \mathbf{H} \cdots A$
01-H1···N1	0.82	1.87	2.580 (3)	145
$N2-H2\cdots O3^{i}$	0.894 (16)	1.991 (15)	2.863 (3)	165
$O3-H3A\cdots O1^{ii}$	0.84 (3)	2.11 (3)	2.895 (3)	155
$O3-H3B\cdots O2$	0.84(2)	2.00(2)	2.829 (3)	167 (3)
$C4-H4\cdots N3^{iii}$	0.93	2.49	3.394 (3)	165
$C7 - H7 \cdots O3^{i}$	0.93	2.56	3.320 (3)	140
$C13-H13\cdots O3^{i}$	0.93	2.47	3.364 (3)	162

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





The packing of (I), viewed along the a axis. Dashed lines indicate intermolecular hydrogen bonds.

The N-H and O-H atoms H2, H3A and H3B were located in a difference Fourier map and refined isotropically, with the N-H distance restrained to 0.90 (1) Å, the O-H distances restrained to 0.85 (1) Å and the H···H distance restrained to 1.37 (2) Å. The other H atoms were placed in idealized positions and constrained to ride on their parent atoms, with an O-H distance of 0.82 Å and C-H distances of 0.93 Å, and with $U_{iso}(H) = 1.2U_{eq}(C)$ and $1.5U_{eq}(O)$.

Data collection: *SMART* (Bruker, 2002); cell refinement: *SAINT* (Bruker, 2002); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997*a*); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997*a*); molecular graphics: *SHELXTL* (Sheldrick, 1997*b*); software used to prepare material for publication: *SHELXL97*.

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