## organic papers

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#### **Key indicators**

Single-crystal X-ray study T = 298 KMean  $\sigma(\text{C}-\text{C}) = 0.007 \text{ Å}$  R factor = 0.038 wR factor = 0.093 Data-to-parameter ratio = 17.5

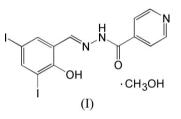
For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e.

# 2'-(2-Hydroxy-3,5-diiodobenzylidene)isonicotinohydrazide methanol solvate

In the title compound,  $C_{13}H_9I_2N_3O_2$ ·CH<sub>3</sub>OH, the Schiff base molecule displays a *trans* configuration with respect to the C—N double bond. The dihedral angle between the benzene and pyridine rings is 24.6 (2)°. The crystal structure is stabilized by intermolecular N-H···O, O-H···N and C-H···O hydrogen bonds, forming a network.

## Comment

Compounds derived from the condensation reaction of aromatic carbaldehydes with hydrazides exhibit a wide range of biological activities and applications (Tarafder *et al.*, 2002; Cukurovali *et al.*, 2002; Ali *et al.*, 2002). The crystal structure of 2'-(2-hydroxy-3,5-diiodobenzylidene)isonicotinohydrazide methanol solvate, (I), is reported here.



The asymmetric unit of (I) comprises a Schiff base molecule and a methanol solvent molecule (Fig. 1), in which the bond lengths and bond angles are within normal ranges (Allen *et al.*, 1987) and comparable with those observed in similar compounds (Qiu *et al.*, 2006; Yang & Guo, 2006; Yang, 2006). The C7=M1 bond length of 1.275 (6) Å conforms to the value for a double bond, and is comparable with that in other Schiff bases (Qian *et al.*, 2006; Zhao, 2006). The C8-M2 bond length of 1.362 (6) Å is intermediate between those typical for the corresponding single and double bonds, suggesting some degree of delocalization in the acetohydrazide system. The molecule displays a *trans* configuration about the C=N and C-N bonds. The dihedral angle between the benzene ring and the pyridine ring is 24.6 (2)°.

The molecular structure is stabilized by an intramolecular  $O-H \cdots N$  hydrogen bond (Table 1). In the crystal structure, molecules are linked through intermolecular  $O-H \cdots N$ ,  $N-H \cdots O$  and  $C-H \cdots O$  hydrogen bonds (Table 1), forming a network (Fig. 2).

## Experimental

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3,5-Diiodosalicylaldehyde (0.1 mmol, 36.9 mg) and pyridine-4carboxylic acid hydrazide (0.1 mmol, 13.7 mg) were dissolved in methanol (10 ml). The mixture was stirred at room temperature for Received 12 January 2007 Accepted 17 January 2007

4887 measured reflections 3490 independent reflections 2579 reflections with  $I > 2\sigma(I)$ 

 $w = 1/[\sigma^2(F_0^2) + (0.0347P)^2]$ 

where  $P = (F_0^2 + 2F_c^2)/3$ 

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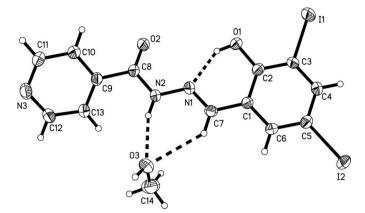
+ 1.1556P]

 $(\Delta/\sigma)_{\rm max} = 0.001$ 

 $\Delta \rho_{\rm max} = 0.96 \ {\rm e} \ {\rm \AA}$ 

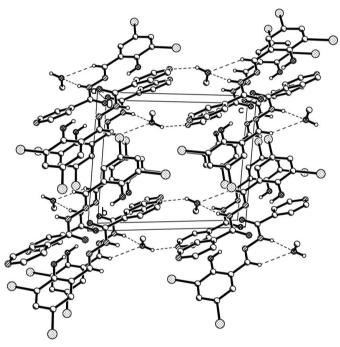
 $\Delta \rho_{\rm min} = -1.09 \text{ e } \text{\AA}^{-3}$ 

 $R_{\rm int} = 0.015$  $\theta_{\rm max} = 26.5^{\circ}$ 



#### Figure 1

The molecular structure of (I), showing 30% probability displacement ellipsoids and the atom-numbering scheme. Hydrogen bonds are shown as dashed lines.



#### Figure 2

The crystal packing of (I), viewed along the *a* axis. Dashed lines indicate hydrogen bonds. H atoms not involved in hydrogen bonding have been omitted.

10 min to give a clear yellow solution. Crystals of (I) were formed by gradual evaporation of the solvent over 10 d at room temperature (yield 82.3%). Analysis found: C 32.17, H 2.59, N 7.87%; calculated for  $C_{14}H_{13}I_2N_3O_3$ : C 32.02, H 2.50, N 8.00%.

### Crystal data

C13H9I2N3O2·CH3OH	V = 860.2 (4) Å <sup>3</sup>
$M_r = 525.07$	Z = 2
Triclinic, $P\overline{1}$	$D_x = 2.027 \text{ Mg m}^{-3}$
a = 8.977 (2) Å	Mo $K\alpha$ radiation
b = 9.114 (3) Å	$\mu = 3.67 \text{ mm}^{-1}$
c = 11.387 (3) Å	T = 298 (2) K
$\alpha = 92.099 \ (2)^{\circ}$	Block, yellow
$\beta = 111.455 \ (3)^{\circ}$	$0.23 \times 0.20 \times 0.18 \text{ mm}$
$\gamma = 95.638 \ (2)^{\circ}$	

#### Data collection

Bruker SMART CCD area-detector diffractometer	
$\omega$ scans	
Absorption correction: multi-scan	
(SADABS; Sheldrick, 1996)	
$T_{\rm min} = 0.486, T_{\rm max} = 0.558$	
(expected range = $0.450-0.517$ )	

## Refinement

Refinement on  $F^2$   $R[F^2 > 2\sigma(F^2)] = 0.038$   $wR(F^2) = 0.093$  S = 1.053490 reflections 200 parameters H-atom parameters constrained

## Table 1

Hydrogen-bond geometry (Å, °).

$D - H \cdots A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdots A$
O1-H1···N1	0.82	1.84	2.555 (5)	146
$N2-H2\cdots O3$	0.86	1.98	2.814 (6)	163
$O3-H3 \cdot \cdot \cdot N3^i$	0.82	2.04	2.786 (6)	152
$C7-H7\cdots O3$	0.93	2.56	3.305 (5)	138

Symmetry code: (i) -x, -y, -z + 1.

H atoms were placed in geometrically idealized positions and allowed to ride on their parent atoms, with O-H = 0.82, N-H = 0.86 and C-H = 0.93-0.96 Å, and with  $U_{iso}(H) = 1.2U_{eq}(C,N)$  or  $1.5U_{eq}$ (methyl C and O). The deepest residual electron density hole is located 0.91 Å from atom I1.

Data collection: *SMART* (Bruker, 1998); cell refinement: *SAINT* (Bruker, 1998); 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: *SHELXTL*.

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