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#### **Key indicators**

Single-crystal X-ray study T = 292 K Mean  $\sigma$ (C–C) = 0.003 Å R factor = 0.037 wR factor = 0.094 Data-to-parameter ratio = 8.5

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

# (E)-N'-(4-Fluorobenzylidene)isonicotinohydrazide monohydrate

In the title molecule,  $C_{13}H_{10}FN_3O \cdot H_2O$ , the hydrazone group adopts a *trans* configuration with respect to the C=N double bond. In the crystal structure, a two-dimensional network is formed *via* intermolecular  $O-H \cdots O$ ,  $N-H \cdots O$  and  $O-H \cdots N$  hydrogen bonds.

### Comment

Schiff base compounds have been of great interest for many years. These compounds play an important role in the development of coordination chemistry related to catalysis and enzymatic reactions (Brunner *et al.*, 1983), magnetism and molecular architectures (Miller & Epstein, 2000).



The molecular structure of the title compound, (I), is shown in Fig. 1. The hydrazone group adopts a *trans* configuration with respect to the C=N double bond. All bond lengths and angles are within normal ranges (Allen *et al.*, 1987). The dihedral angle between the benzene and pyridine rings is  $11.5 (2)^{\circ}$ .

In the crystal structure, a two-dimensional network, in the *ab* plane, is formed *via* intermolecular  $O-H\cdots O$ ,  $N-H\cdots O$  and  $O-H\cdots N$  hydrogen bonds (Table 2 and Fig. 2).

### Experimental

4-Fluorobenzaldehyde (0.2 mmol, 24.8 mg) and isonicotinohydrazide (0.2 mmol, 27.4 mg) were dissolved in methanol (15 ml). The mixture was stirred at room temperature for about 30 min and the resulting solution was set aside for 10 d to allow slow evaporation of the solvent. Large colourless block-shaped crystals of (I) separated from the solution; these were collected and washed three times with methanol.

Crystal data

 $C_{13}H_{10}FN_{3}O \cdot H_{2}O$   $M_{r} = 261.26$ Orthorhombic,  $P2_{1}2_{1}2_{1}$  a = 6.469 (2) Å b = 7.021 (2) Å c = 27.351 (9) Å  $V = 1242.3 (7) \text{ Å}^{3}$ 

Z = 4  $D_x$  = 1.397 Mg m<sup>-3</sup> Mo K $\alpha$  radiation  $\mu$  = 0.11 mm<sup>-1</sup> T = 292 (2) K Block, colourless 0.30 × 0.30 × 0.20 mm Received 24 October 2006 Accepted 27 October 2006

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#### Data collection

Bruker SMART CCD area-detector diffractometer  $\varphi$  and  $\omega$  scans Absorption correction: multi-scan (*SADABS*; Sheldrick, 1996)  $T_{\rm min} = 0.969, T_{\rm max} = 0.979$ 

#### Refinement

Refinement on  $F^2$   $R[F^2 > 2\sigma(F^2)] = 0.037$   $wR(F^2) = 0.094$  S = 0.981562 reflections 184 parameters 5749 measured reflections 1562 independent reflections 1163 reflections with  $I > 2\sigma(I)$  $R_{\text{int}} = 0.029$  $\theta_{\text{max}} = 27.0^{\circ}$ 

H atoms treated by a mixture of independent and constrained refinement  $w = 1/[\sigma^2(F_o^2) + (0.0531P)^2]$ where  $P = (F_o^2 + 2F_c^2)/3$  $(\Delta/\sigma)_{max} < 0.001$  $\Delta\rho_{max} = 0.16$  e Å<sup>-3</sup> $\Delta\rho_{min} = -0.13$  e Å<sup>-3</sup>

## Table 1

Hydrogen-bond geometry (Å,  $^{\circ}$ ).

$D - H \cdot \cdot \cdot A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - H \cdots A$
$N2-H1N\cdotsO1W^{i}$	0.91 (2)	1.95 (2)	2.832 (3)	164 (2)
$O1W - H1WB \cdots O1^{ii}$	0.834 (10)	2.28 (2)	2.949 (2)	138 (3)
O1W−H1WB···N3 <sup>ii</sup>	0.834 (10)	2.54 (2)	3.264 (3)	146 (3)
O1W-H1 $WA$ ···N1 <sup>iii</sup>	0.844 (10)	2.048 (13)	2.863 (3)	162 (2)
	4 4 555		4	

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

In the absence of significant anomalous dispersion effects, Friedel pairs were merged. The H atoms of the water molecules and imine group were located in a difference Fourier map and refined isotropically, with water H atoms refined with the O-H and H···H distances restrained to 0.84 (1) and 1.37 (2) Å, respectively. The other H atoms were placed in geometrically idealized positions, with C-H = 0.93 Å, and constrained to ride on their parent atoms, with  $U_{iso}(H) = 1.2U_{eq}(C)$ 

Data collection: *SMART* (Siemens, 1996); cell refinement: *SAINT* (Siemens, 1996); 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*) and *PLATON* (Spek, 2003); software used to prepare material for publication: *SHELXL97*.



#### Figure 1

The molecular structure of (I), showing 30% probability displacement ellipsoids and the atom-numbering scheme.





Part of the crystal structure of (I). Dashed lines indicate hydrogen bonds.

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