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Key indicators

Single-crystal X-ray study T = 293 K Mean σ (C–C) = 0.003 Å R factor = 0.036 wR factor = 0.098 Data-to-parameter ratio = 14.0

For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e. In the title compound, $C_7H_5N_3OS$, the dihedral angle between the pyridyl and oxadiazole rings is 9.32 (2) Å. In the crystal structure, a combination of intermolecular N-H···N hydrogen bonds and π - π stacking interactions forms a two-

5-(3-Pyridyl)-1,3,4-oxadiazole-2-thione

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Comment

dimensional network.

The design and assembly of supramolecular architectures with variable cavities or channels are currently of great interest owing to their novel topologies and potential applications (Wang et al., 2005a,b; Wang, Tong et al., 2005). The geometry of the organic ligands has a great effect on the structural frameworks of such coordination polymers and hence much effort has been devoted to the modification of the building blocks and the control of the assembled motifs for the required products through the selection of different organic ligands. Previous studies have shown that mostly symmetric rigid bridging ligands are used in the construction of versatile networks and other porous coordination polymers (Robin & Fromm, 2006). However, there are fewer studies on the coordination chemistry of unsymmetric organic ligands (Wang, Tang & Qin, 2006; Wang, Tang, Wu et al., 2006). We have recently discussed some coordination complexes of 5-(4pyridyl)-1,3,4-oxadiazole-2-thione (Wang & Tang 2006*a*,*b*) and reported the crystal structure of the free ligand (Wang, Tang, Ma et al., 2006). To further expand this area of research, we have synthesized the title compound, (I), by the reaction of nicotinoylhydrazide and carbon disulfide in the presence of KOH in ethanol and its crystal structure is presented here.



The molecular structure of (I) is shown in Fig. 1. All bond lengths and angles are within the expected ranges (Allen *et al.*, 1987). In the crystal structure, one-dimensional chains are formed *via* intermolecular $N-H\cdots N$ hydrogen bonds (Table 2). In addition, $\pi-\pi$ stacking interactions $[Cg1\cdots Cg2(1 + x, y, z) = 3.591$ (2) Å, where Cg1 and Cg2 are the centroids of the ring atoms O1/C6/N2/N3/C7 and N1/C1– C5, respectively] connect one-dimensional chains into a twodimensional network (Fig. 2).

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Experimental

The title compound was synthesized according to a literature method (Wang, Tang, Ma et al., 2006). Pale-yellow plate-shaped single crystals suitable for X-ray diffraction were obtained from an ethanol solution of (I). Analysis found (%): C 46.80, H 2.82, N 23.40; requires (%): C 46.92, H 2.81, N 23.45.

Z = 4

 $D_x = 1.503 \text{ Mg m}^{-3}$

Mo $K\alpha$ radiation

Plate, pale yellow

 $0.30\,\times\,0.20\,\times\,0.10$ mm

4047 measured reflections

1804 independent reflections

1371 reflections with $I > 2\sigma(I)$

 $\mu = 0.36 \text{ mm}^{-1}$

T = 293 (2) K

 $R_{\rm int} = 0.021$

 $\theta_{\rm max} = 27.6^\circ$

Crystal data

C7H5N3OS $M_{\rm m} = 179.20$ Monoclinic, $P2_1/c$ a = 5.1351 (2) Å b = 18.1940 (7) Å c = 8.5727 (4) Å $\beta = 98.694 \ (3)^{\circ}$ V = 791.73 (6) Å³

Data collection

Bruker SMART APEX CCD diffractometer φ and ω scans Absorption correction: multi-scan (SADABS; Sheldrick, 1996) $T_{\min} = 0.900, \ T_{\max} = 0.965$

Refinement

Refinement on F^2	$w = 1/[\sigma^2(F_0^2) + (0.0475P)^2]$
$R[F^2 > 2\sigma(F^2)] = 0.036$	+ 0.162P]
$wR(F^2) = 0.098$	where $P = (F_0^2 + 2F_c^2)/3$
S = 1.00	$(\Delta/\sigma)_{\rm max} = 0.001$
1804 reflections	$\Delta \rho_{\rm max} = 0.21 \ {\rm e} \ {\rm \AA}^{-3}$
129 parameters	$\Delta \rho_{\rm min} = -0.25 \text{ e} \text{ \AA}^{-3}$
All H-atom parameters refined	

Table 1

Hydrogen-bond geometry (Å, °).

$D - H \cdot \cdot \cdot A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - H \cdots A$
$N3-H1A\cdots N1^{i}$	0.94 (2)	1.86 (2)	2.800 (2)	172 (2)
Symmetry code: (i) r	$-1 - v + \frac{3}{2} = +$	1		

Symmetry code: (i) $x - 1, -y + \frac{3}{2}, z + \frac{1}{2}$.

All H atoms were refined independently with isotropic displacement parameters, giving N-H = 0.94 (2) Å and C-H = 0.924 (18)-0.95 (2) Å.

Data collection: SMART (Bruker, 2001); cell refinement: SAINT (Bruker, 2001); data reduction: SAINT; program(s) used to solve structure: SHELXS97 (Sheldrick, 1997); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: SHELXTL (Sheldrick, 1999); software used to prepare material for publication: SHELXTL.

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Figure 1

The molecular structure of (I), showing displacement ellipsoids drawn at the 30% probability level.



Figure 2

Part of the crystal structure of (I), showing hydrogen bonding interactions aa dashed lines.

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