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## Structure Reports

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## Key indicators

Single-crystal X-ray study
$T=293 \mathrm{~K}$
Mean $\sigma(\mathrm{C}-\mathrm{C})=0.003 \AA$
$R$ factor $=0.036$
$w R$ 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.

[^0]
## 5-(3-Pyridyl)-1,3,4-oxadiazole-2-thione

In the title compound, $\mathrm{C}_{7} \mathrm{H}_{5} \mathrm{~N}_{3} \mathrm{OS}$, the dihedral angle between the pyridyl and oxadiazole rings is 9.32 (2) $\AA$. In the crystal structure, a combination of intermolecular $\mathrm{N}-\mathrm{H} \cdots \mathrm{N}$ hydrogen bonds and $\pi-\pi$ stacking interactions forms a twodimensional network.

## Comment

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-(4-pyridyl)-1,3,4-oxadiazole-2-thione (Wang \& Tang 2006a,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.

(I)

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 $\mathrm{N}-\mathrm{H} \cdots \mathrm{N}$ hydrogen bonds (Table 2). In addition, $\pi-\pi$ stacking interactions $[\operatorname{Cg} 1 \cdots \operatorname{Cg} 2(1+x, y, z)=3.591$ (2) $\AA$, where $C g 1$ and $C g 2$ are the centroids of the ring atoms $\mathrm{O} 1 / \mathrm{C} 6 / \mathrm{N} 2 / \mathrm{N} 3 / \mathrm{C} 7$ and $\mathrm{N} 1 / \mathrm{C} 1-$ 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 .

## Crystal data

$\mathrm{C}_{7} \mathrm{H}_{5} \mathrm{~N}_{3} \mathrm{OS}$
$M_{r}=179.20$
Monoclinic, $P 2_{\mathrm{b}} / c$
$a=5.1351(2) \AA$
$b=18.1940(7) \AA$
$c=8.5727(4) \AA$
$\beta=98.694(3){ }^{\circ}{ }^{\circ}$
$V=791.73(6) \AA^{3}$

## Data collection

Bruker SMART APEX CCD diffractometer
$\varphi$ and $\omega$ scans
Absorption correction: multi-scan (SADABS; Sheldrick, 1996)

$$
\begin{aligned}
& Z=4 \\
& D_{x}=1.503 \mathrm{Mg} \mathrm{~m}^{-3}
\end{aligned}
$$

Mo $K \alpha$ radiation
$a=5.1351$ (2) A
$\mu=0.36 \mathrm{~mm}^{-1}$
$T=293$ (2) K
Plate, pale yellow
$0.30 \times 0.20 \times 0.10 \mathrm{~mm}$
$T_{\text {min }}=0.900, T_{\text {max }}=0.965$

## Refinement

Refinement on $F^{2}$

$$
\begin{aligned}
& w=1 /\left[\sigma^{2}\left(F_{\mathrm{o}}{ }^{2}\right)+(0.0475 P)^{2}\right. \\
& +0.162 P \text { ] } \\
& \text { where } P=\left(F_{\mathrm{o}}^{2}+2 F_{\mathrm{c}}{ }^{2}\right) / 3 \\
& (\Delta / \sigma)_{\max }=0.001 \text { 。 } \\
& \Delta \rho_{\text {max }}=0.21 \mathrm{e}^{-3} \\
& \Delta \rho_{\min }=-0.25 \mathrm{e}^{-3}
\end{aligned}
$$

$w R\left(F^{2}\right)=0.098$
$S=1.00$
1804 reflections
129 parameters
All H -atom parameters refined

4047 measured reflections 1804 independent reflections 1371 reflections with $I>2 \sigma(I)$
$R_{\text {int }}=0.021$
$\theta_{\text {max }}=27.6^{\circ}$

Table 1
Hydrogen-bond geometry ( $\AA,{ }^{\circ}$ ).

| $D-\mathrm{H} \cdots A$ | $D-\mathrm{H}$ | $\mathrm{H} \cdots A$ | $D \cdots A$ | $D-\mathrm{H} \cdots A$ |
| :--- | :--- | :--- | :--- | :--- |
| $\mathrm{~N} 3-\mathrm{H} 1 A \cdots \mathrm{~N} 1^{\mathrm{i}}$ | $0.94(2)$ | $1.86(2)$ | $2.800(2)$ | $172(2)$ |

Symmetry code: (i) $x-1,-y+\frac{3}{2}, z+\frac{1}{2}$.
All H atoms were refined independently with isotropic displacement parameters, giving $\mathrm{N}-\mathrm{H}=0.94$ (2) $\AA$ and $\mathrm{C}-\mathrm{H}=0.924$ (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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