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#### Key indicators

Single-crystal X-ray study T = 93 K Mean  $\sigma$ (N–C) = 0.002 Å Disorder in main residue R factor = 0.034 wR factor = 0.095 Data-to-parameter ratio = 14.2

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# N'-(2-Methoxythiobenzoyl)pyridine-2-carbohydrazide

In the crystal structure of the title compound,  $C_{14}H_{13}N_3O_2S$ , intermolecular N-H····O (H···O = 2.03 Å) and C-H···S (H···S = 2.77 Å) hydrogen bonds link molecules into a two-dimensional framework.

## Comment

In recent years, there has been considerable interest in metal complexes of nitrogen-sulfur donor ligands, such as thiohydrazides, whose properties can be altered by introducing different organic substituents. These ligands can coordinate to metal ions to give structures with different geometries and properties (Singh *et al.*, 1988; Shrivastav *et al.*, 2002; Liu *et al.*, 1992). As part of our ongoing research on these ligands, we report here the crystal structure of the title compound, (I), a new thiohydrazide.



The molecular structure of (I), together with the atomlabeling scheme, is shown in Fig. 1. The molecule is not exactly planar, but can be divided into three planar fragments, viz. an o-methoxyphenyl ring, a carbonothioyl carbohydrazide unit and a pyridine ring. The peripheral rings are disordered over two orientations, with occupancies of 0.556 (6) and 0.444 (6) for the pyridine ring and 0.519 (3) and 0.481 (3) for the omethoxyphenyl ring (Fig. 2). The carbohydrazide unit is almost coplanar with the pyridine ring [dihedral angles = 3.26 (3) and 4.44 (6) $^{\circ}$  for the major and minor components, respectively]. The o-methoxyphenyl ring plane is twisted away from the carbonothioyl carbohydrazide plane [60.2 (2) and 55.9 (2)° for the major and minor components, respectively]. This twisting may be attributed to an intermolecular C2- $H2A \cdots O2$  hydrogen bond. The dihedral angles between the o-methoxyphenyl ring and the pyridine ring are 61.3 (3) and  $60.0 (3)^{\circ}$  for the major and minor components, respectively. Hydrazinic atoms H2B and H3B are trans to each other, as are the C7-S and C6-O1 groups [torsion angles N3-N2-C6-O1 and N2-N3-C7-S = -1.8 (2) and 1.2 (2)°, respectively]. In addition, the C–S bond distance of 1.669 (1) Å agrees well with equivalent bonds in similar structures, being intermediate between 1.82 Å for a C–S single bond and 1.56 Å for a C=S



#### Figure 1

The molecular structure of (I), showing the atom-numbering scheme. Displacement ellipsoids are drawn at the 20% probability level. H atoms are shown as small spheres of arbitrary radii. Only one disorder component is shown. The dashed line indicates a hydrogen bond.



Figure 2

The molecular structure of (I), showing the disordered rings. H atoms have been omitted for clarity.

double bond (Wu et al., 2000). The corresponding C7-N3 bond distance of 1.3279 (16) Å is also intermediate between 1.47 Å for a C–N single bond and 1.29 Å for a double bond (Boga et al., 1999). The N2-N3 distance of 1.3767 (13) Å also shows partial double-bond character, suggesting extensive delocalization in the compound. The zigzag conformation of (I) makes atoms S, N2 and N1 potential donor sites for coordination with metals. Intermolecular N-H···O, C- $H \cdots O$  and  $C - H \cdots S$  hydrogen bonds link the molecules together in the solid state (Table 2 and Fig. 2).

# **Experimental**

Compound (I) was synthesized by the reaction of 2-methoxyphenylcarbonothioylthioacetic acid and picolinic acid hydrazide in an alkaline medium. The phenylcarbonothioylthioacetic acid was prepared according to the literature method of Jensen & Pedersen (1961). Solutions of 2-methoxyphenylcarbonothioylthioacetic acid (2.42 g, 10 mmol) and picolinic acid hydrazide (1.36 g, 10 mmol) in 1 N NaOH were mixed together and allowed to stand for 2 h. The resulting filtered solution was acidified with acetic acid (20% v/v). The precipitated product was filtered off, washed twice with portions of cold water and dried in vacuo. White single crystals of (I) (m.p. 431 K) suitable for X-ray analysis were obtained by slow evaporation of a methanol-diethyl ether (50:50) solution over a period of 4 d (yield 1.4 g, 68%). IR (v, cm<sup>-1</sup>): 3248 and 3184 (-NH), 1666 (-C=O), 912 (-C=S); <sup>1</sup>H NMR (DMSO-d<sub>6</sub>, TMS): 3.88 (s, 3H, -OCH<sub>3</sub>), 7.03-7.46 (m, 4H, aromatic), 7.98-8.75 (m, 4H, pyridine), 11.79 and 13.20 (s, 1H from -NH, D<sub>2</sub>O exchangeable); <sup>13</sup>C NMR (DMSO-d<sub>6</sub>, TMS): 149.08 (C1), 127.53 (C2), 138.23 (C3), 120.33 (C4), 147.77 (C5), 159.33



#### Figure 3

A projection of the crystal packing of (I) along the b axis. Only one disorder component is shown. The dashed lines indicate hydrogen bonds.

(C6), 184.91 (C7), 122.32 (C8), 131.65 (C9), 127.05 (C10), 111.96 (C11), 131.32 (C12), 155.13 (C13), 56.00 (C14).

## Crystal data

$C_{14}H_{13}N_3O_2S$	$D_x = 1.412 \text{ Mg m}^{-3}$
$M_r = 287.33$	Mo $K\alpha$ radiation
Monoclinic, $P2_1/n$	Cell parameters from 5777
a = 14.608 (2)  Å	reflections
b = 6.7090 (11)  Å	$\theta = 3.0-28.3^{\circ}$
c = 14.975 (3) Å	$\mu = 0.24 \text{ mm}^{-1}$
$\beta = 112.946 \ (3)^{\circ}$	T = 93 (2) K
V = 1351.5 (4) Å <sup>3</sup>	Chunk, colorless
Z = 4	$0.80 \times 0.55 \times 0.45 \ \text{mm}$

#### Data collection

Bruker SMART CCD area-detector	3221 independent reflections
diffractometer	2639 reflections with $I > 2\sigma(I)$
$\varphi$ and $\omega$ scans	$R_{\rm int} = 0.031$
Absorption correction: multi-scan	$\theta_{\rm max} = 28.3^{\circ}$
(SADABS; Bruker, 1999)	$h = -19 \rightarrow 18$
$T_{\min} = 0.710, \ T_{\max} = 0.896$	$k = -8 \rightarrow 8$
10072 measured reflections	$l = -19 \rightarrow 17$

## Refinement

Refinement on $F^2$	$w = 1/[\sigma^2(F_0^2) + (0.0487P)^2]$
$R[F^2 > 2\sigma(F^2)] = 0.034$	+ 0.3326P]
$vR(F^2) = 0.095$	where $P = (F_0^2 + 2F_c^2)/3$
S = 1.05	$(\Delta/\sigma)_{\rm max} = 0.001$
221 reflections	$\Delta \rho_{\rm max} = 0.26 \text{ e } \text{\AA}^{-3}$
27 parameters	$\Delta \rho_{\rm min} = -0.29 \text{ e } \text{\AA}^{-3}$
H-atom parameters constrained	

# Table 1

Selected geometric parameters (Å,  $^{\circ}$ ).

S-C7	1.6690 (12)	N3-C7	1.3279 (16)
O1-C6	1.2299 (17)	C5A - C6	1.587 (10)
N2-C6	1.3357 (17)	C5B-C6	1.446 (8)
N2-N3	1.3767 (13)		
C6-N2-N3	121.41 (10)	C7-N3-N2	119.80 (10)

Table 2			
Hydrogen-bond geometry	(Å,	°).	

$D - H \cdots A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdot \cdot \cdot A$
$N2-H2A\cdots N1A$	0.88	2.17	2.639 (11)	113
$N3-H3A\cdotsO1^{i}$	0.88	2.03	2.8524 (14)	155
$C2A - H2AA \cdots O2A^{ii}$	0.95	2.35	3.152 (13)	141
$C2B - H2BA \cdots O2A^{ii}$	0.95	2.45	3.268 (13)	145
$C12A - H12A \cdots S^{iii}$	0.95	2.77	3.584 (6)	144
Symmetry codes: (i)	-x + 1, -	-y, -z; (ii)	) $-x + \frac{3}{2}, y - \frac{1}{2},$	$-z + \frac{1}{2};$ (iii)

 $x - \frac{1}{2}, -y + \frac{1}{2}, z - \frac{1}{2}.$ 

The peripheral rings of (I) are disordered over two orientations, as described in the *Comment*. The two components were restrained to have similar geometric and displacement parameters. All H atoms were initially located in a difference Fourier map. The methyl H atoms were then constrained to an ideal geometry, with C-H = 0.98 Å and  $U_{iso}(H) = 1.5U_{eq}(C)$ , but each group was allowed to rotate freely about its C–C bond. The amine H atoms, as well as those attached to C atoms, were placed in geometrically idealized positions and constrained to ride on their parent atoms, with N–H = 0.88 Å, C–H = 0.95–1.00 Å and  $U_{iso}(H) = 1.2U_{eq}(C,N)$ .

Data collection: *SMART* (Bruker, 1998); cell refinement: *SAINT-Plus* (Bruker, 1999); data reduction: *SAINT-Plus*; program(s) used to

solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *SHELXTL* (Bruker, 1999); software used to prepare material for publication: *SHELXTL*.

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