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FangFang Jian* and Ying Li

New Materials & Function Coordination Chemistry Laboratory, Qingdao University of Science & Technology, Qingdao 266042, People's Republic of China

Correspondence e-mail: ffj2003@163169.net

Key indicators

Single-crystal X-ray study T = 296 KMean $\sigma(\text{C}-\text{C}) = 0.004 \text{ Å}$ R factor = 0.056 wR factor = 0.180 Data-to-parameter ratio = 17.4

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

1-(Furan-2-ylmethylene)-4-phenylthiosemicarbazide

In the molecule of the title compound, $C_{12}H_{11}N_3OS$, the dihedral angle between the planar rings is 59.4 (2)°. The structure is stabilized by intramolecular $N-H\cdots N$ and intermolecular $N-H\cdots S$ hydrogen bonds. Intermolecular $N-H\cdots S$ hydrogen bonds link the independent molecules into dimers.

Comment

Thiosemicarbazide is interesting because of the formation of complexes with biological activities (Shen *et al.*, 1998). Some substituted thiourea derivatives have shown interesting biological effects, including anti-HIV properties (Mao *et al.*, 1999), and thiourea derivatives have also been successfully screened for various biological actions (Antholine & Taketa, 1982). As a ligand with potential S- and N-atom donors, thiosemicarbazide is interesting because of the structural chemistry of its multifunctional coordination modes (*N*-monodentate, *S*-monodentate or *N*:*S*-bidentate). In order to investigate further this kind of ligand, we synthesized the title compound, (I), and describe its structure here.



In the molecule of (I) (Fig. 1), the bond lengths and angles are in normal ranges (Allen *et al.*, 1987; Ji *et al.*, 2002). The dihedral angle between the planar rings is $59.4 (2)^{\circ}$.



Figure 1

The molecular structure of (I) with the atom-numbering scheme. Displacement ellipsoids are drawn at the 30% probability level. The intramolecular hydrogen bond is shown as a dashed line.

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The crystal structure of (I) is stabilized by intramolecular $N-H\cdots N$ and intermolecular $N-H\cdots S$ hydrogen bonds (Table 1). Intermolecular $N-H\cdots S$ hydrogen bonds link the independent molecules into dimers (Fig. 2). Dipole–dipole and van der Waals interactions are also effective in the molecular packing in the crystal structure.

Experimental

The title compound was prepared by the reaction of hydrazine (1.0 g, 20 mmol) and furfural (1.9 g, 20 mmol) with phenyl isothiocyanate (2.7 g, 20 mmol). Single crystals suitable for X-ray measurements were obtained by recrystallization from an acetone solution at room temperature (yield 3.9 g, 79.4%; m.p. 402–404 K).

V = 611.6 (2) Å³

 $D_x = 1.332 \text{ Mg m}^{-3}$

 $0.35 \times 0.25 \times 0.25 \mbox{ mm}$

3 standard reflections

every 100 reflections

intensity decay: none

 $w = 1/[\sigma^2(F_o^2) + (0.1127P)^2]$

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

+ 0.1125P]

 $\Delta \rho_{\rm max} = 0.40 \text{ e} \text{ Å}^{-3}$

 $\Delta \rho_{\rm min} = -0.52 \ {\rm e} \ {\rm \AA}^{-3}$

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

1882 reflections with $I > 2\sigma(I)$

Mo $K\alpha$ radiation

 $\mu = 0.25 \text{ mm}^{-1}$

T = 296 (2) K

Block, brown

 $R_{\rm int} = 0.010$

 $\theta_{\rm max} = 27.0^{\circ}$

Z = 2

Crystal data

 $\begin{array}{l} C_{12}H_{11}N_3OS\\ M_r = 245.31\\ \text{Triclinic, } P\overline{1}\\ a = 5.8810 \ (12) \ \text{\AA}\\ b = 9.878 \ (2) \ \text{\AA}\\ c = 11.186 \ (2) \ \text{\AA}\\ \alpha = 71.45 \ (3)^{\circ}\\ \beta = 83.18 \ (3)^{\circ}\\ \gamma = 88.95 \ (3)^{\circ} \end{array}$

Data collection

Enraf–Nonius CAD-4 diffractometer ω scans Absorption correction: none 2889 measured reflections 2675 independent reflections

Refinement

Refinement on F^2 $R[F^2 > 2\sigma(F^2)] = 0.056$ $wR(F^2) = 0.180$ S = 1.062675 reflections 154 parameters H-atom parameters constrained

Table 1

Hydrogen-bond geometry (Å, °).

$D - H \cdot \cdot \cdot A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdot \cdot \cdot A$
$N1-H1A\cdots N3$	0.86	2.19	2.607 (3)	109
$N2-H2A\cdots S1^{i}$	0.86	2.54	3.382 (2)	166

Symmetry code: (i) -x, -y + 2, -z + 1.







H atoms were positioned geometrically, with N-H = 0.86 Å and C-H = 0.93 Å, and constrained to ride on their parent atoms, with $U_{\rm iso}({\rm H}) = 1.2U_{\rm eq}({\rm C,N}).$

Data collection: *CAD-4 Software* (Enraf–Nonius, 1989); cell refinement: *CAD-4 Software*; data reduction: *NRCVAX* (Gabe *et al.*, 1989); program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *SHELXTL/PC* (Sheldrick, 1990); software used to prepare material for publication: *WinGX* (Farrugia, 1999).

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