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2'-(2-Thienylmethylidene)-2-hydroxybenzohydrazide

Fu-You Pan and Jian-Guo Yang*

Department of Chemistry, Taizhou University, Taizhou 317000, People's Republic of China

Correspondence e-mail: yjg@tzc.edu.cn

Key indicators

Single-crystal X-ray study T = 293 KMean $\sigma(C-C) = 0.003 \text{ Å}$ R factor = 0.052wR factor = 0.157 Data-to-parameter ratio = 14.7

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

The title compound, C₁₂H₁₀N₂O₂S, was synthesized by the reaction of 2-hydroxybenzoylhydrazine with 2-thiophenecarboxaldehyde in ethanol. The crystal structure involves intermolecular O-H···O and intramolecular N-H···O hydrogen bonds.

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Comment

Some benzoylhydrazone compounds possess bacteriostatic activity. This type of compound has wide application in the treatment of tuberculosis and also exhibits fungicidal activity (Edwards et al., 1975). Furthermore, the hydrazonecarbonyl is a structural motif showing bioactivity (Zhi et al., 2003). In order to search for more effective antibacterial medicines, we have synthesized the title compound, (I).

The molecule is essentially planar, with an r.m.s. deviation of 0.0048 Å. An intermolecular O−H···O hydrogen bond is observed linking the hydroxyl H atom with the carbonyl group of an adjacent molecule. In addition, an intramolecular O-H···N hydrogen bond linking the amide NH group with the hydroxyl group, forming a six-membered ring (Fig. 2), is also found.

Experimental

2-Hydroxybenzoylhydrazine (0.02 mol, 3.04 g) was dissolved in anhydrous ethanol (50 ml) at room temperature. 2-Thiophenecarbox-

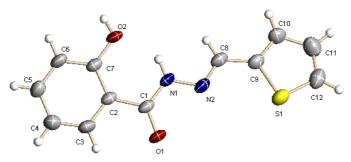


Figure 1 The structure of (I), showing the atomic numbering. Displacement ellipsoids are drawn at the 30% probability level.

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aldehyde (0.02 mol, 1.88 g) was added and the mixture refluxed for 2 h. The precipitate was collected by filtration and washed with ethanol. The product was recrystallized from ethanol and dried under reduced pressure, giving the title compound. The compound (2.5 mmol, 0.62 g) was dissolved in dimethylformamide (30 ml) and kept at room temperature for 30 d, after which colorless blockshaped single crystals formed and were collected and washed with distilled water.

Crystal data

$C_{12}H_{10}N_2O_2S$	$D_x = 1.460 \text{ Mg m}^{-3}$
$M_r = 246.28$	Mo $K\alpha$ radiation
Monoclinic, $P2_1/n$	Cell parameters from 2410
a = 4.9785 (6) Å	reflections
b = 21.127 (2) Å	$\theta = 5.4-54.4^{\circ}$
c = 10.9004 (13) Å	$\mu = 0.28 \text{ mm}^{-1}$
$\beta = 102.209 (2)^{\circ}$	T = 293 (2) K
$V = 1120.6 (2) \text{ Å}^3$	Block, colorless
Z = 4	$0.51 \times 0.26 \times 0.17 \text{ mm}$

Data collection

Bruker SMART APEX area-	2442 independent reflections
detector diffractometer	1945 reflections with $I > 2\sigma(I)$
φ and ω scans	$R_{\rm int} = 0.073$
Absorption correction: multi-scan	$\theta_{\rm max} = 27.0^{\circ}$
(SADABS; Bruker, 2002)	$h = -6 \rightarrow 6$
$T_{\min} = 0.871, T_{\max} = 0.955$	$k = -18 \rightarrow 26$
6565 measured reflections	$l = -13 \rightarrow 12$

Refinement

Refinement	
Refinement on F^2	H atoms treated by a mixture of
$R[F^2 > 2\sigma(F^2)] = 0.052$	independent and constrained
$wR(F^2) = 0.157$	refinement
S = 1.04	$w = 1/[\sigma^2(F_o^2) + (0.0957P)^2]$
2442 reflections	where $P = (F_o^2 + 2F_c^2)/3$
166 parameters	$(\Delta/\sigma)_{\rm max} < 0.001$
	$\Delta \rho_{\text{max}} = 0.36 \text{ e Å}^{-3}$
	$\Delta \rho_{\min} = -0.36 \text{ e Å}^{-3}$

Table 1 Selected bond lengths (Å).

S1-C12	1.682(2)	N1-C1	1.344 (3)
S1-C9	1.702 (2)	N1-N2	1.368 (2)
O1-C1	1.228 (2)	N2-C8	1.268 (3)

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

$D-H\cdots A$	D-H	$H \cdot \cdot \cdot A$	$D \cdot \cdot \cdot A$	$D-\mathrm{H}\cdots A$
$ \begin{array}{c} O2-H2\cdotsO1^{i}\\ N1-H1\cdotsO2 \end{array} $	0.781 (17)	1.918 (18)	2.6681 (19)	161 (3)
	0.80 (2)	2.01 (2)	2.625 (2)	134.5 (19)

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

All H atoms (except H1, H2 and H8) were positioned geometrically and allowed to ride on their parent atoms at distances of

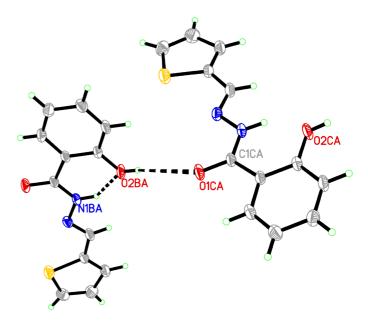


Figure 2 Intermolecular and intramolecular hydrogen bonds (dashed lines).

0.93 Å (C—H), with $U_{\rm iso}({\rm H})=1.2U_{\rm eq}({\rm parent\ atom})$. Atoms H1, H2 and H8 were located in a difference Fourier map and their parameters were refined. The N1—H1 distance is 0.80 (2) Å, the O2—H2 distance is 0.781 (17) Å and the C8—H8 distance is 0.89 (3) Å.

Data collection: *SMART* (Bruker, 2002); cell refinement: *SAINT* (Bruker, 2002); data reduction: *SAINT* and *SHELXTL* (Bruker, 2002); program(s) used to solve structure: *SHELXS*97 (Sheldrick, 1997); program(s) used to refine structure: *SHELXL*97 (Sheldrick, 1997); molecular graphics: *SHELXTL*; software used to prepare material for publication: *SHELXTL*.

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