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Ming-Li Liu, Jian-Min Dou,* Da-Qi Wang and Da-Cheng Li

School of Chemistry and Chemical Engineering, Liaocheng University, Liaocheng 252059, People's Republic of China

Correspondence e-mail: jmdou@lctu.edu.cn

Key indicators

Single-crystal X-ray study T = 298 KMean σ (C–C) = 0.004 Å R factor = 0.043 wR factor = 0.129 Data-to-parameter ratio = 10.0

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

2-Hydroxy-3-methoxybenzaldehyde (pyridine-2-carbonyl)hydrazone

The approximately planar molecule of the title compound, $C_{14}H_{13}N_3O_3$, is in an *E* configuration with respect to the C=N bond, with a C-C=N-N torsion angle of -177.3 (2)°. There is an intramolecular hydrogen bond involving the hydrazide N atom and the hydroxyl O atom on the benzene ring. Intermolecular C-H···O weak interactions are also formed between two adjacent methoxy groups, linking two molecules into a coplanar dimer. These dimers are further assembled into a zigzag framework along the [010] axis, using intermolecular short C-H···O and long N-H···O hydrogen bonds, resulting in a two-dimensional supramolecular crystal structure.

Comment

The chemistry of aroylhydrazones continues to attract much attention due to their coordination ability to metal ions (Singh *et al.*, 1982; Salem, 1998) and their biological activity (Singh *et al.*, 1982; Carcelli *et al.*, 1995). In particular, aroylhydrazones containing a pyridine ring present high sensitivity and selectivity to some metals and have therefore been applied as analytical reagents (Singh *et al.*, 1982; Iki *et al.*, 1994, 1997; Babaiah *et al.*, 1996). Such compounds as N'-salicylidene-3-pyridinecarbohydrazide, 2-[1-(salicyloylhydrazono)ethyl]pyridinium chloride dihydrate and 2-(2-hydroxybenzylidene)-1-(2-picoloyl)hydrazine have been synthesized and characterized by X-ray diffraction (Abboud *et al.*, 1997; Wang *et al.*, 1998; Galić *et al.*, 2001). We now report on another compound, (I), whose structure was determined by X-ray diffraction.



The molecular structure of the title compound is shown in Fig. 1. This compound contains two aromatic rings linked through a monoacylhydrazone group. An *E* configuration with respect to the C=N bond is shown by the molecule, with a C-C=N-N torsion angle of -177.3 (2)°. Bond lengths C1=O1 and C7=N2 are 1.221 (3) and 1.281 (3) Å, respectively, in agreement with double-bond character, whereas bonds C1-N1, C8-C7, C9-O2 and C10-O3 are typical single bonds (Table 1). Bond lengths observed in (I) are in agreement with values found in the related compounds, *N'*-salicylidene-3-pyridinecarbohydrazide (Galić *et al.*, 2001) and isonicotinic acid(2-hydroxy-3-methoxybenzilidene)hydrazide (Yu *et al.*,

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2322 independent reflections 1261 reflections with $I > 2\sigma(I)$

 $w = 1/[\sigma^2(F_0^2) + (0.0558P)^2]$

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

+ 0.0418P]

 $\Delta \rho_{\rm min} = -0.16 \text{ e } \text{\AA}^{-3}$

 $(\Delta/\sigma)_{\rm max} < 0.001$ $\Delta \rho_{\rm max} = 0.19 \text{ e } \text{\AA}^{-3}$

 $R_{\rm int}=0.052$ $\theta_{\rm max} = 25.0^{\circ}$ $h = -19 \rightarrow 19$ $k = -7 \rightarrow 8$ $l = -13 \rightarrow 12$



Figure 1

The molecular structure of (I). Displacement ellipsoids are drawn at the 30% probability level.



Figure 2 The crystal packing of (I).

2005). These values suggest that the aroylhydrazone moiety in (I) exists in the ketoamino form.

The molecule of the title compound is almost planar. Three relevant dihedral angles indicative of a slight deviation from planarity are: $7.60 (12)^{\circ}$ between the benzene and pyridine rings; $5.06 (13)^{\circ}$ between the benzene ring and the central hydrazone linkage (O1/C1/N1/N2/C7); 5.25 (12)° between the pyridine ring and the central hydrazone linkage. The intramolecular hydrogen bond involving the hydroxy group bonded to the benzene ring and atom N2 of the hydrazone chain stabilizes this configuration. Moreover, a search of the Cambridge Structural Database (CSD; Version 5.18; Allen, 2002) for unsubstituted salicylideneamines in the enolimino form revealed 165 fragments in 123 structures, with structures invariably almost planar and including an intramolecular O- $H \cdots N$ hydrogen bond.

Atoms C14 and O3 of the methoxy group in one molecule act as donor and acceptor to form a weak intermolecular hydrogen bond (Table 2). As a result, a stable six-membered ring is built with two symmetry-related molecules linked into a planar dimer. These dimers are further hydrogen bonded: each dimer provides one carbonyl O atom, which interacts with NH and CH groups of the hydrazone group of a neighbouring molecule, giving a lzigzag framework along the [010] axis (Fig. 2). It is worth noting that the $C7 \cdots O1^{ii}$ separation is significantly shorter than the N1···O1ⁱⁱ separation [symmetry code: (ii) $x, -y + \frac{1}{2}, z + \frac{1}{2}$].

Experimental

A solution of pyridinehydrazide (10.0 mmol, 1.371 g) in methanol (15 ml) was mixed with o-vanillin (14.0 mmol, 2.134 g) in methanol (25 ml) and stirred for 2 h at ca 323 K. The yellow precipitate was filtered off and crystallized from methanol–DMF (1:1 v/v), yielding crystals suitable for X-ray diffraction (m.p. 448-450 K). FT-IR spectrum: 3432, 3272, 2923, 2850, 1669, 1607, 775, 729, 707 cm⁻¹.

Crystal data

C ₁₄ H ₁₃ N ₃ O ₃	$D_x = 1.358 \text{ Mg m}^{-3}$
$M_r = 271.27$	Mo $K\alpha$ radiation
Monoclinic, $P2_1/c$	Cell parameters from 1287
a = 16.534(5)Å	reflections
b = 7.538 (2) Å	$\theta = 2.6-20.6^{\circ}$
c = 11.228 (3) Å	$\mu = 0.10 \text{ mm}^{-1}$
$\beta = 108.554 \ (5)^{\circ}$	T = 298 (2) K
$V = 1326.6 (7) \text{ Å}^3$	Block, yellow
Z = 4	$0.43 \times 0.38 \times 0.21 \text{ mm}$

Data collection

Refinement

Refinement on F^2 $R[F^2 > 2\sigma(F^2)] = 0.044$ wR(F²) = 0.129 S = 1.022322 reflections 233 parameters All H-atom parameters refined

Table 1

Selected geometric parameters (Å, °).

N1-C1	1.345 (3)	O2-C9	1.353 (3)
N1-N2	1.378 (3)	O3-C10	1.371 (3)
N2-C7	1.281 (3)	O3-C14	1.420 (3)
O1-C1	1.221 (3)		
C1-N1-N2	120.3 (2)	O1-C1-N1	124.0 (2)
C7-N2-N1	116.7 (2)	O1-C1-C2	122.7 (3)
C10-O3-C14	117.9 (3)	N1-C1-C2	113.3 (2)
N2-N1-C1-C2	-177.2 (2)	N1-N2-C7-C8	-177.3 (2)

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

$D - H \cdot \cdot \cdot A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdot \cdot \cdot A$
$C7 - H6 \cdots O1^{i}$	1.00(3)	2.29(3)	3.213 (3)	153 (2)
$O2 - H7 \cdots N2$	0.89(2)	1.81(2)	2.618 (2)	149 (3)
$N1-H1\cdotsO1^{i}$	0.84 (3)	2.66 (3)	3.402 (3)	148 (2)
C14-H11···O3 ⁱⁱ	1.00 (3)	2.41 (3)	3.264 (5)	142 (2)

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

All H atoms were located in a difference Fourier map and freely refined isotropically [C-H = 0.91 (3)–1.02(3) Å].

Data collection: *SMART* (Siemens, 1996); cell refinement: *SAINT* (Siemens, 1996); 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, 1997); software used to prepare material for publication: *SHELXTL*.

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