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## Key indicators

Single-crystal X-ray study

T = 294 K

Mean  $\sigma(\text{C}-\text{C}) = 0.002 \text{ \AA}$ 

R factor = 0.037

wR factor = 0.103

Data-to-parameter ratio = 13.7

For details of how these key indicators were  
automatically derived from the article, see  
<http://journals.iucr.org/e>.*N,N'*-Bis(3-hydroxy-4-methoxybenzylidene)-  
hydrazine

The title compound,  $\text{C}_{16}\text{H}_{16}\text{N}_2\text{O}_4$ , was synthesized by the reaction of 3-hydroxy-4-methoxybenzaldehyde with hydrazine hydrate. The molecule possesses a crystallographically imposed centre of symmetry. An intramolecular  $\text{O}-\text{H}\cdots\text{O}$  hydrogen bond promotes planarity of the molecular backbone. In the crystal structure, intermolecular  $\text{O}-\text{H}\cdots\text{N}$  and  $\text{C}-\text{H}\cdots\text{O}$  hydrogen bonds stabilize the crystal packing.

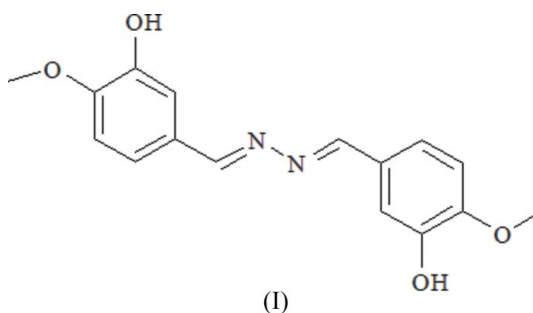
Received 1 September 2005

Accepted 21 September 2005

Online 28 September 2005

## Comment

Recently, a number of azine compounds containing both a diimine linkage and N—N bonding have been investigated in terms of their crystallography and coordination chemistry (Zheng *et al.*, 2005; Kundu *et al.*, 2005; Kessler & Euler, 1999; Armstrong *et al.*, 1998; Xu *et al.*, 1997). We report here the crystal structure of the title compound, (I), where two 3-hydroxy-4-methoxybenzylidene units are directly linked through the imine N atoms.



The molecule of (I) possesses a crystallographically imposed centre of symmetry at the mid-point of the N—N bond (Fig. 1). The N—N bond length of 1.412 (2) Å (Table 1) is somewhat longer than that observed in related azine compounds (Xu *et al.*, 2005; Liu *et al.*, 2004; Şengül *et al.*, 2004).

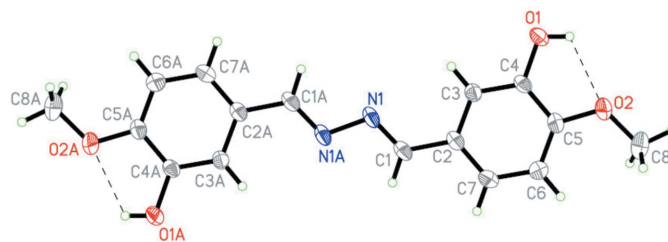
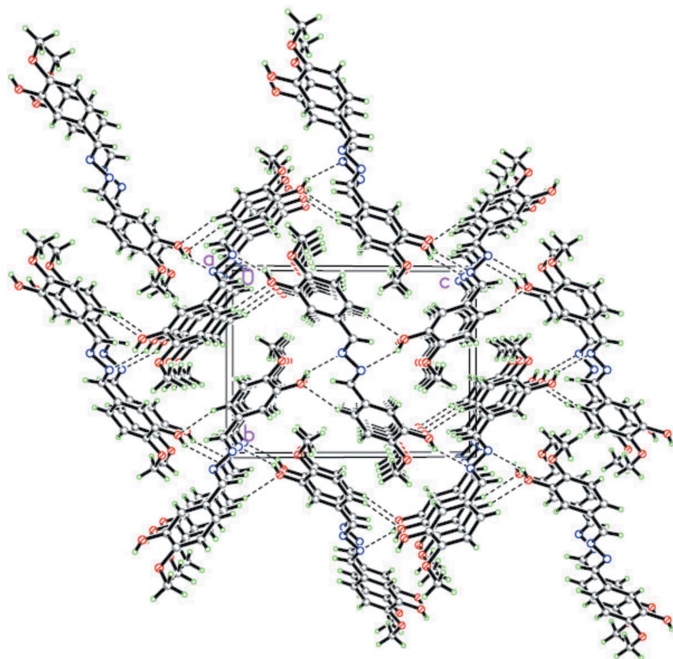


Figure 1

A view of (I), showing the intramolecular hydrogen bonds (dashed lines) and with displacement ellipsoids at the 30% probability level. The suffix A corresponds to symmetry operation  $(-1 - x, 1 - y, 1 - z)$ .



**Figure 2**  
The crystal packing of (I), viewed approximately down the *a* axis, showing the intermolecular hydrogen-bonded (dashed lines) extended network.

The C=N–N angle [ $112.82(15)^\circ$ ] is similar to that in *N,N'*-bis(4-chlorobenzylidene)hydrazine (Zheng *et al.*, 2005) but significantly smaller than the ideal  $sp^2$  value of  $120^\circ$ , as a consequence of repulsion between the nitrogen lone pairs and the adjacent C=N bond. An intramolecular O–H...O hydrogen bond (Table 2) provides planarity of the molecular backbone, with an r.m.s deviation of  $0.034(6)$  Å.

In the crystal structure, intermolecular O–H...N and C–H...O hydrogen bonds (Table 2) stabilize the crystal packing (Fig. 2).

## Experimental

The title compound was synthesized by the reaction of 3-hydroxy-4-methoxybenzaldehyde with hydrazine hydrate in refluxing ethanol (Liu *et al.*, 2004). Single crystals suitable for X-ray analysis were obtained by slow evaporation of a tetrahydrofuran solution.

### Crystal data

$C_{16}H_{16}N_2O_4$	$D_x = 1.339$ Mg m $^{-3}$
$M_r = 300.31$	Mo $K\alpha$ radiation
Monoclinic, $P2_1/c$	Cell parameters from 1595 reflections
$a = 5.6858(12)$ Å	$\theta = 2.6$ – $26.1^\circ$
$b = 10.018(2)$ Å	$\mu = 0.10$ mm $^{-1}$
$c = 13.902(3)$ Å	$T = 294(2)$ K
$\beta = 109.841(4)^\circ$	Block, light yellow
$V = 744.8(3)$ Å $^3$	$0.30 \times 0.22 \times 0.16$ mm
$Z = 2$	

### Data collection

Bruker SMART CCD area-detector diffractometer	1523 independent reflections
$\varphi$ and $\omega$ scans	1065 reflections with $I > 2\sigma(I)$
Absorption correction: multi-scan (SADABS; Sheldrick, 1996)	$R_{int} = 0.026$
$T_{min} = 0.946$ , $T_{max} = 0.980$	$\theta_{max} = 26.4^\circ$
4096 measured reflections	$h = -5 \rightarrow 7$
	$k = -12 \rightarrow 10$
	$l = -17 \rightarrow 15$

### Refinement

Refinement on $F^2$	$w = 1/[\sigma^2(F_o^2) + (0.0456P)^2 + 0.1273P]$
$R[F^2 > 2\sigma(F^2)] = 0.037$	where $P = (F_o^2 + 2F_c^2)/3$
$wR(F^2) = 0.103$	$(\Delta/\sigma)_{max} < 0.001$
$S = 1.03$	$\Delta\rho_{max} = 0.12$ e Å $^{-3}$
1523 reflections	$\Delta\rho_{min} = -0.18$ e Å $^{-3}$
111 parameters	
H atoms treated by a mixture of independent and constrained refinement	

**Table 1**

Selected geometric parameters (Å, °).

O1–C4	1.3659 (17)	N1–C1	1.269 (2)
O2–C5	1.3631 (17)	N1–N1 <sup>i</sup>	1.413 (2)
O2–C8	1.417 (2)		
C5–O2–C8	118.80 (13)	N1–C1–C2	122.33 (13)
C1–N1–N1 <sup>i</sup>	112.82 (15)	O1–C4–C3	119.14 (14)
N1 <sup>i</sup> –N1–C1–C2	177.44 (14)	N1–C1–C2–C3	–9.5 (2)
N1–C1–C2–C7	171.79 (14)		

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

**Table 2**

Hydrogen-bond geometry (Å, °).

<i>D</i> –H... <i>A</i>	<i>D</i> –H	H... <i>A</i>	<i>D</i> ... <i>A</i>	<i>D</i> –H... <i>A</i>
O1–H1...O2	0.86 (1)	2.20 (2)	2.6405 (17)	112 (2)
O1–H1...N1 <sup>ii</sup>	0.86 (1)	2.18 (2)	2.9365 (16)	147 (2)
C7–H7...O1 <sup>iii</sup>	0.93	2.49	3.391 (2)	164

Symmetry codes: (ii)  $-x, y + \frac{1}{2}, -z + \frac{3}{2}$ ; (iii)  $x, -y + \frac{3}{2}, z - \frac{1}{2}$ .

All H atoms were located in a difference Fourier map. C-bound H atoms were refined as riding, with C–H =  $0.93$ – $0.96$  Å and  $U_{iso}(H) = 1.2U_{eq}(C)$ . Atom H1 was refined isotropically, with O1–H1 =  $0.85(1)$  Å.

Data collection: SMART (Bruker, 1997); cell refinement: SAINT (Bruker, 1997); data reduction: SAINT; program(s) used to solve structure: SHELXS97 (Sheldrick, 1997); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: SHELXTL (Bruker, 1997); software used to prepare material for publication: SHELXTL.

The authors gratefully acknowledge financial support from the Foundation for Excellent Young Teachers of Jiangxi Science and Technology Normal University.

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