

(E)-1-(4-Ethoxy-3-methoxybenzylidene)-2-(4-nitrophenyl)hydrazine**Jun Shi†**

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Key indicatorsSingle-crystal X-ray study
 $T = 294$ K
Mean $\sigma(\text{C}-\text{C}) = 0.008$ Å
 R factor = 0.051
 wR factor = 0.146
Data-to-parameter ratio = 6.7For details of how these key indicators were automatically derived from the article, see <http://journals.iucr.org/e>.

The title compound, $\text{C}_{16}\text{H}_{17}\text{N}_3\text{O}_4$, crystallizes with two molecules in the asymmetric unit. Each independent molecule is nearly planar. Two bifurcated intermolecular $\text{N}-\text{H}\cdots(\text{O},\text{O})$ hydrogen-bond systems help to consolidate the crystal packing.

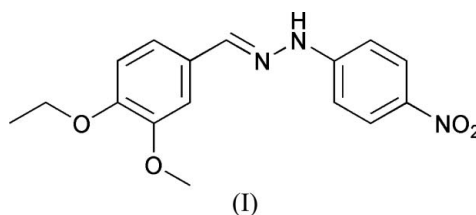
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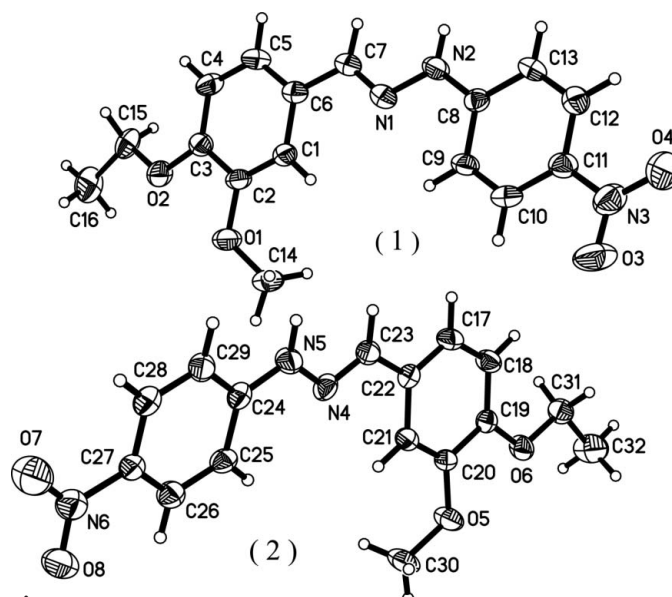
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Comment

Metal complexes based on Schiff bases have attracted much attention because of their biological activity (Kahwa *et al.*, 1986). Consequently, a large number of Schiff base derivatives have been synthesized and employed to develop protein and enzyme mimics, such as models to mimic hydrolase in the hydrolysis of *p*-nitrophenyl picolinate (Li *et al.*, 2005).



As part of an investigation of their crystal structures, which will provide useful information for the coordination properties of Schiff bases functioning as ligands, in the present study we report the synthesis and molecular structure of the nitrophenylhydrazine Schiff base derivative (I) (Fig. 1).

**Figure 1**

The structure of the asymmetric unit of (I), with displacement ellipsoids for non-H atoms drawn at the 30% probability level.

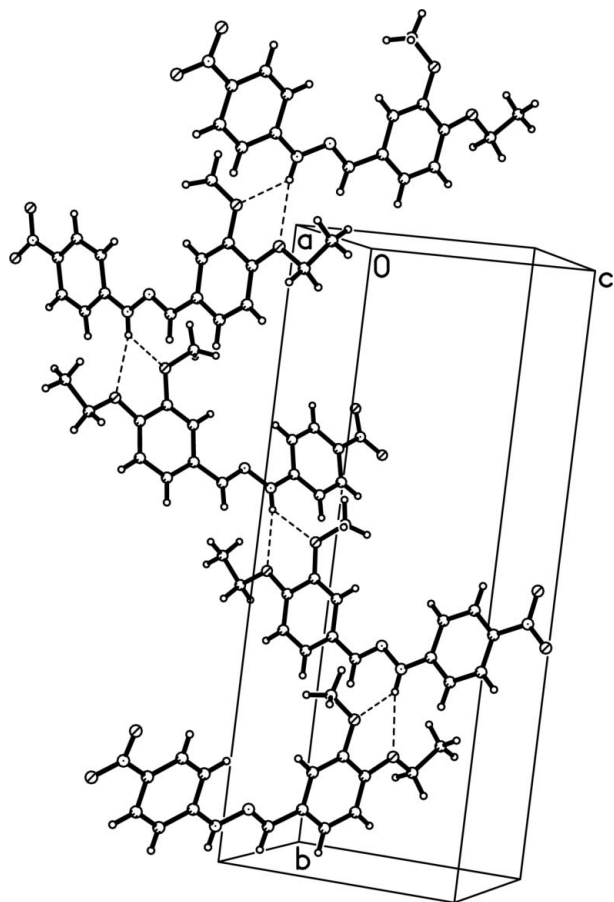


Figure 2
Packing diagram for (I), with H bonds drawn as dashed lines.

The asymmetric unit of (I) consists of two independent molecules, which are quite similar to each other. In molecule 1, the vanillin group (C1–C7/O1/O2) is planar, with an r.m.s. deviation, δ , from the mean plane of 0.029 Å, and it makes a dihedral angle of 11.29 (18)° with the phenylhydrazine residue (C8–C13/N1/N2). In molecule 2, the vanillin group (C17–C23/O5/O6) is also planar, with $\delta = 0.027$ Å, and it makes a dihedral angle of 10.87 (13)° with the phenylhydrazine residue (C24–C29/N4/N5). The nitro group and its attached aromatic ring are not coplanar, with dihedral angles of 7.23 (9)° (for O3/N3/O4) and 4.61 (6)° (for O7/N6/O8). Furthermore, the phenylhydrazine residues of the two independent molecules are almost perpendicular to each other, with a dihedral angle of 86.36 (9)°, while the dihedral angle between the two vanillin groups is 77.10 (9)°. All bond lengths and angles are within normal ranges (Allen *et al.*, 1987).

Two bifurcated intermolecular N–H···(O,O) hydrogen-bond systems are found in (I) (Table 1 and Fig. 2), which help to consolidate the crystal packing. The hydrogen bonds link adjacent molecules, forming infinite chains.

Experimental

An anhydrous ethanol solution of 4-ethoxy-3-methoxybenzaldehyde (1.80 g, 10 mmol) was added to an anhydrous ethanol solution of 1-(4-nitrophenyl)hydrazine (1.53 g, 10 mmol) and the mixture stirred at

350 K for 5 h under nitrogen. A yellow precipitate appeared which was isolated, recrystallized from acetonitrile, and then dried in a vacuum to give the pure compound in 83% yield. Yellow single crystals of (I) suitable for X-ray analysis were obtained by slow evaporation of an acetonitrile solution.

Crystal data

$C_{16}H_{17}N_3O_4$
 $M_r = 315.33$
 Monoclinic, $P2_1$
 $a = 7.589$ (2) Å
 $b = 23.491$ (7) Å
 $c = 8.929$ (2) Å
 $\beta = 96.287$ (5)°
 $V = 1582.2$ (7) Å³
 $Z = 4$

$D_x = 1.324$ Mg m⁻³
 Mo $K\alpha$ radiation
 Cell parameters from 1909 reflections
 $\theta = 2.8$ – 22.1 °
 $\mu = 0.10$ mm⁻¹
 $T = 294$ (2) K
 Block, yellow
 $0.22 \times 0.20 \times 0.18$ mm

Data collection

Bruker SMART APEX CCD area-detector diffractometer
 φ and ω scans
 Absorption correction: multi-scan (SADABS; Sheldrick, 1996)
 $T_{\min} = 0.962$, $T_{\max} = 0.983$
 7919 measured reflections

2820 independent reflections
 1741 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.044$
 $\theta_{\text{max}} = 25.0$ °
 $h = -8 \rightarrow 6$
 $k = -27 \rightarrow 27$
 $l = -10 \rightarrow 10$

Refinement

Refinement on F^2
 $R[F^2 > 2\sigma(F^2)] = 0.051$
 $wR(F^2) = 0.146$
 $S = 1.14$
 2820 reflections
 419 parameters

H-atom parameters constrained
 $w = 1/[\sigma^2(F_o^2) + (0.072P)^2]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\text{max}} = 0.021$
 $\Delta\rho_{\text{max}} = 0.31$ e Å⁻³
 $\Delta\rho_{\text{min}} = -0.21$ e Å⁻³

Table 1

Hydrogen-bond geometry (Å, °).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
N2–H2···O5 ⁱ	0.86	2.39	3.042 (6)	133
N2–H2···O6 ⁱ	0.86	2.36	3.179 (6)	160
N5–H5···O1 ⁱⁱ	0.86	2.37	2.949 (6)	125
N5–H5···O2 ⁱⁱ	0.86	2.36	3.198 (7)	166

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

The H atoms were included in calculated positions and refined using a riding-model approximation. Constrained C–H and N–H bond lengths and $U_{\text{iso}}(\text{H})$ parameters: 0.93 Å and $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$ for aromatic CH; 0.97 Å and $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$ for methylene CH₂; 0.96 Å and $U_{\text{iso}}(\text{H}) = 1.5U_{\text{eq}}(\text{C})$ for methyl CH₃; 0.86 Å and $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{N})$ for imino NH. In the absence of significant anomalous dispersion effects, Friedel pairs were averaged.

Data collection: SMART (Bruker, 1999); cell refinement: SAINT (Bruker, 1999); data reduction: SAINT; program(s) used to solve structure: SHELXS97 (Sheldrick, 1997a); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997a); molecular graphics: SHELXTL (Sheldrick, 1997b); software used to prepare material for publication: SHELXTL.

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