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Key indicators

Single-crystal X-ray study
 $T = 298$ K
Mean $\sigma(\text{C}-\text{C}) = 0.011$ Å
 R factor = 0.144
 wR factor = 0.278
Data-to-parameter ratio = 13.7For details of how these key indicators were
automatically derived from the article, see
<http://journals.iucr.org/e>.

2-Methoxybenzaldehyde isonicotinoylhydrazone

The title molecule, $\text{C}_{14}\text{H}_{13}\text{N}_3\text{O}_2$, adopts a *trans* configuration with respect to the $\text{C}=\text{N}$ double bond. There are two molecules in the asymmetric unit. The dihedral angles between the two rings are 39.1 (4) and 19.7 (4)°. In the crystal structure, intermolecular $\text{N}-\text{H}\cdots\text{O}$ and $\text{N}-\text{H}\cdots\text{N}$ hydrogen bonds generate a network structure.

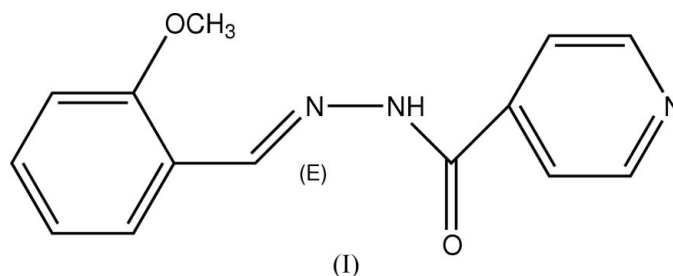
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Comment

The background to this study is described in the previous paper (Xie *et al.*, 2006). As an extension of work on the structural characterization of hydrazone Schiff base compounds, we report here the crystal structure of (I), a new isonicotinohydrazone with 2-methoxybenzaldehyde.



In the title compound, (I), which crystallizes with two unique molecules in the asymmetric unit (Fig. 1), the $\text{C}-\text{N}$ bonds in the hydrazone units are characteristically short (Table 1) because of conjugation effects. All other bond lengths are within normal ranges (Allen *et al.*, 1987). The dihedral angles between the benzene and pyridine rings are 39.1 (4)° ($\text{C}1/\text{C}2/\text{C}3/\text{N}1/\text{C}4/\text{C}5$ with $\text{C}8/\text{C}9/\text{C}10/\text{C}11/\text{C}12/\text{C}13$) and 19.7 (4)° ($\text{C}15/\text{C}16/\text{C}17/\text{N}4/\text{C}18/\text{C}19$ with $\text{C}22/\text{C}23/\text{C}24/\text{C}25/\text{C}26/\text{C}27$); these are slightly larger than normal (Fun *et al.*, 1997) due to the steric effect of the $\text{C}13$ and $\text{C}27$ methoxy substituents. The crystal structure is stabilized by intermolecular $\text{N}-\text{H}\cdots\text{O}$ and $\text{N}-\text{H}\cdots\text{N}$ hydrogen bonds (Table 2 and Fig. 2)

Experimental

2-Methoxybenzaldehyde (0.2 mmol, 27.2 mg) and isonicotinohydrazide (0.2 mmol, 27.4 mg) were dissolved in methanol (10 ml). The mixture was stirred at room temperature for about 10 min. to give a clear yellow solution. The solution was set aside for 8 d to allow slow evaporation of the solvent. Large colourless plate-shaped crystals separated from the solution; these were collected and washed three times with water.

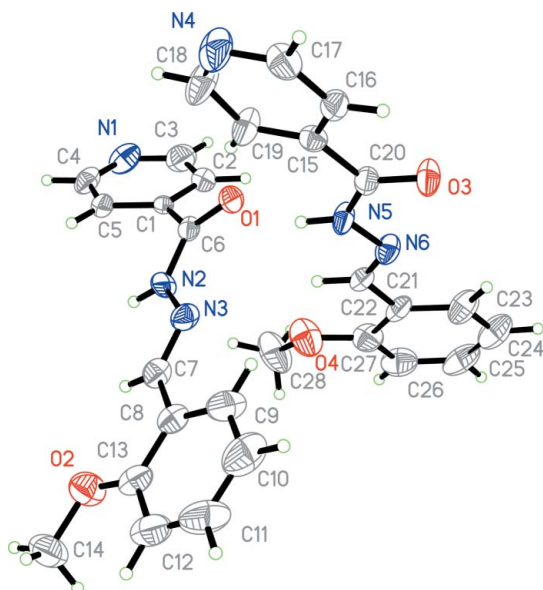


Figure 1
The structure of the asymmetric unit of (I), showing 30% probability displacement ellipsoids and the atom-numbering scheme.

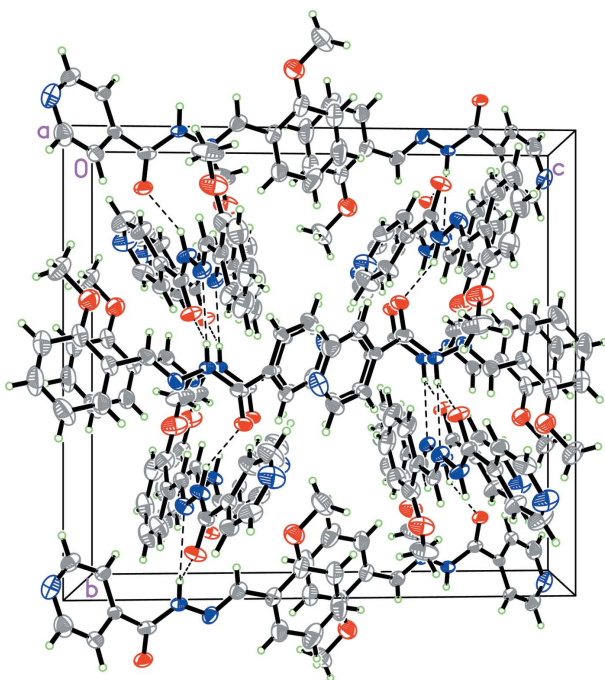


Figure 2
The crystal packing of (I), viewed along the *a* axis. Dashed lines indicate intermolecular hydrogen bonds.

Crystal data

$C_{14}H_{13}N_3O_2$
 $M_r = 255.27$
 Monoclinic, $P2_1/c$
 $a = 9.766$ (2) Å
 $b = 15.935$ (4) Å
 $c = 17.428$ (4) Å
 $\beta = 95.404$ (4)°
 $V = 2700.1$ (10) Å³
 $Z = 8$

$D_x = 1.256$ Mg m⁻³
 Mo $K\alpha$ radiation
 Cell parameters from 1047 reflections
 $\theta = 4.9$ – 35.8 °
 $\mu = 0.09$ mm⁻¹
 $T = 298$ (2) K
 Plate, colourless
 $0.40 \times 0.20 \times 0.10$ mm

Data collection

Bruker SMART APEX area-detector diffractometer
 φ and ω scans
 Absorption correction: multi-scan (SADABS; Bruker, 2002)
 $T_{\min} = 0.966$, $T_{\max} = 0.991$
 13018 measured reflections

4742 independent reflections
 2711 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.090$
 $\theta_{\text{max}} = 25.0$ °
 $h = -11 \rightarrow 11$
 $k = -17 \rightarrow 18$
 $l = -20 \rightarrow 18$

Refinement

Refinement on F^2
 $R[F^2 > 2\sigma(F^2)] = 0.144$
 $wR(F^2) = 0.278$
 $S = 1.24$
 4742 reflections
 345 parameters
 H-atom parameters constrained

$w = 1/[\sigma^2(F_o^2) + (0.0652P)^2 + 1.9118P]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\text{max}} = 0.007$
 $\Delta\rho_{\text{max}} = 0.27$ e Å⁻³
 $\Delta\rho_{\text{min}} = -0.19$ e Å⁻³

Table 1

Selected geometric parameters (Å, °).

O1—C6	1.221 (6)	N5—C20	1.327 (7)
O3—C20	1.216 (6)	N5—N6	1.364 (6)
N2—C6	1.340 (8)	N6—C21	1.269 (7)
N2—N3	1.366 (7)		
C6—N2—N3	120.8 (5)	C21—N6—N5	115.1 (5)
C7—N3—N2	113.8 (5)	O1—C6—N2	124.1 (7)
C20—N5—N6	119.6 (5)	O3—C20—N5	122.9 (6)
C6—N2—N3—C7	−178.0 (6)	C7—C8—C13—O2	−0.3 (10)
C20—N5—N6—C21	−178.3 (6)	C21—C22—C27—O4	−1.0 (10)

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>
N2—H2A...O3 ⁱ	0.86	2.09	2.825 (6)	143
N2—H2A...N6 ⁱ	0.86	2.55	3.294 (7)	146
N5—H5A...O1	0.86	2.07	2.800 (6)	142

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

All H atoms were placed in geometrically idealized positions [$C-H = 0.93$ (aromatic H atoms) or 0.96 Å (methyl H atoms); $N-H = 0.86$ Å], and refined as riding atoms, with $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{aromatic C, N})$ or $1.5U_{\text{eq}}(\text{methyl C})$.

Data collection: SMART (Bruker, 2002); cell refinement: SAINT (Bruker, 2002); 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, 2000); software used to prepare material for publication: SHELXL97.

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