

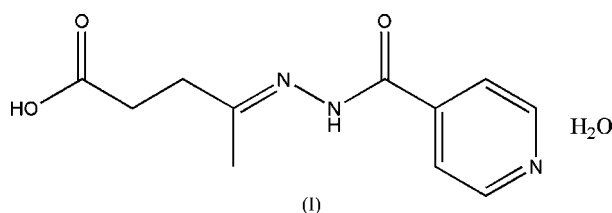
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Key indicators

Single-crystal X-ray study
 $T = 293$ K
Mean $\sigma(\text{C}-\text{C}) = 0.004$ Å
 R factor = 0.080
 wR factor = 0.185
Data-to-parameter ratio = 13.9For details of how these key indicators were
automatically derived from the article, see
<http://journals.iucr.org/e>.**(E)-4-(Isonicotinoylhydrazono)pentanoic
acid monohydrate**The title compound, $\text{C}_{11}\text{H}_{13}\text{N}_3\text{O}_3 \cdot \text{H}_2\text{O}$, was synthesized by the reaction of acetopropanoic acid and isonicotinoylhydrazide in ethanol. A three-dimensional network of intermolecular $\text{O}-\text{H} \cdots \text{O}$, $\text{O}-\text{H} \cdots \text{N}$ and $\text{N}-\text{H} \cdots \text{O}$ hydrogen bonds stabilizes the crystal structure.Received 28 February 2006
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Comment

Hydrazine and its derivatives exhibit biological and pharmacological activities (Vicini *et al.*, 2002; Maccari *et al.*, 2005). These compounds are also used as synthetic intermediates (Rollas *et al.*, 2002). The title compound, (I), was synthesized in our laboratory by the reaction of acetopropanoic acid and isonicotinoylhydrazide in an ethanol medium. We present here its crystal structure (Fig. 1).The bond lengths (Table 1) and angles show normal values. The uncoordinated water molecule plays an important role in the formation of a three-dimensional network of intermolecular $\text{O}-\text{H} \cdots \text{O}$ and $\text{N}-\text{H} \cdots \text{O}$ hydrogen bonds (Table 2), which stabilize the crystal structure (Fig. 2). $\text{O}-\text{H} \cdots \text{N}$ interactions are also present.

Experimental

A mixture of acetopropanoic acid (1.16 mg, 10 mmol) and isonicotinoylhydrazide (1.37 mg, 10 mmol) was refluxed in ethanol for 3 h. After cooling, the mixture was filtered and dried. The title compound was recrystallized from a mixed solvent of methanol and water (1:5) in 85% yield (200 mg). Block-shaped colourless single crystals suitable for X-ray diffraction were obtained. Analysis found (%): C 52.03, H 5.96, N 16.66; $\text{C}_{11}\text{H}_{15}\text{N}_3\text{O}_4$ requires (%): C 52.17, H 5.97, N 16.59.

Crystal data

 $\text{C}_{11}\text{H}_{13}\text{N}_3\text{O}_3 \cdot \text{H}_2\text{O}$
 $M_r = 253.26$
Monoclinic, $P2_1/c$
 $a = 9.7834$ (14) Å
 $b = 12.4798$ (17) Å
 $c = 10.0924$ (15) Å
 $\beta = 92.447$ (3)°
 $V = 1231.1$ (3) Å³
 $Z = 4$ $D_x = 1.366$ Mg m⁻³
Mo $K\alpha$ radiation
Cell parameters from 6637
reflections
 $\theta = 2-52^\circ$
 $\mu = 0.11$ mm⁻¹
 $T = 293$ (2) K
Block, colourless
 $0.30 \times 0.30 \times 0.20$ mm

Data collection

Bruker SMART CCD area-detector diffractometer
 φ and ω scans
 Absorption correction: multi-scan (SADABS; Sheldrick, 1996)
 $T_{\min} = 0.969$, $T_{\max} = 0.979$
 6637 measured reflections

2425 independent reflections
 1674 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.038$
 $\theta_{\text{max}} = 26.0^\circ$
 $h = -5 \rightarrow 12$
 $k = -15 \rightarrow 14$
 $l = -12 \rightarrow 12$

Refinement

Refinement on F^2
 $R[F^2 > 2\sigma(F^2)] = 0.080$
 $wR(F^2) = 0.185$
 $S = 1.23$
 2425 reflections
 175 parameters
 H atoms treated by a mixture of independent and constrained refinement

$w = 1/[\sigma^2(F_o^2) + (0.0716P)^2 + 0.1162P]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\text{max}} < 0.001$
 $\Delta\rho_{\text{max}} = 0.28 \text{ e } \text{Å}^{-3}$
 $\Delta\rho_{\text{min}} = -0.29 \text{ e } \text{Å}^{-3}$

Table 1

Selected bond lengths (Å).

O1—C6	1.224 (3)	N1—C5	1.331 (4)
O2—C11	1.321 (4)	N2—C6	1.342 (3)
O3—C11	1.204 (3)	N2—N3	1.396 (3)
N1—C1	1.325 (4)	N3—C7	1.273 (3)

Table 2

Hydrogen-bond geometry (Å, °).

$D-H \cdots A$	$D-H$	$H \cdots A$	$D \cdots A$	$D-H \cdots A$
O1W—H1WA \cdots O1 ⁱ	0.86 (5)	1.92 (5)	2.769 (4)	171 (4)
N2—H2 \cdots O1W	0.86	2.09	2.869 (3)	151
O1W—H1WB \cdots O2 ⁱⁱ	0.86 (5)	2.04 (5)	2.887 (3)	168 (4)
O2—H2C \cdots N1 ⁱⁱⁱ	0.85 (4)	1.81 (4)	2.647 (4)	170 (4)

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

All H atoms were initially located in a difference Fourier map. The O-bound H atoms were refined isotropically. All other H atoms were placed in geometrically idealized positions and refined as riding, with $N-H = 0.86 \text{ Å}$, $C-H = 0.93-0.97 \text{ Å}$ and $U_{\text{iso}}(H) = 1.2U_{\text{eq}}$ of the parent atom.

Data collection: SMART (Bruker, 2001); cell refinement: SAINT (Bruker, 2001); data reduction: SAINT; program(s) used to solve structure: SHELXS97 (Sheldrick, 1997); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics:

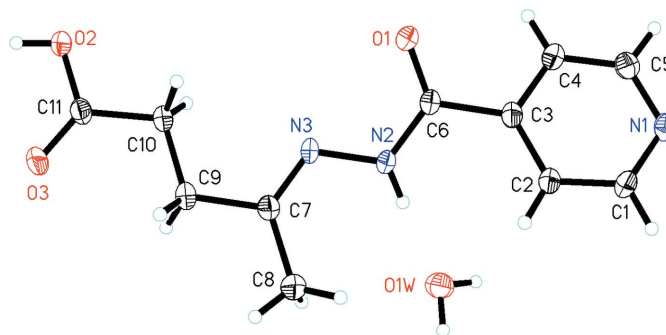


Figure 1

View of (I), showing displacement ellipsoids drawn at the 30% probability level. H atoms are represented by circles of arbitrary size.

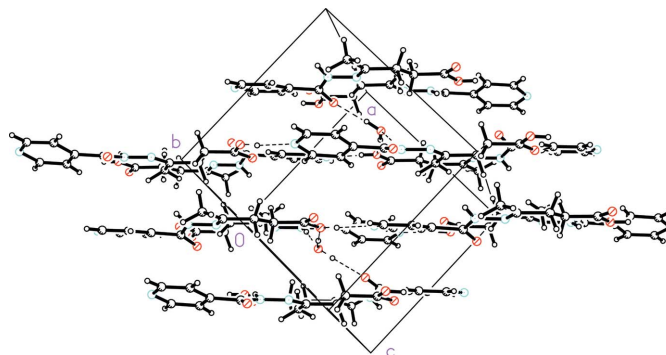


Figure 2

Packing diagram, showing the intermolecular hydrogen bonds as dashed lines.

SHELXTL (Sheldrick, 1999); software used to prepare material for publication: SHELXTL.

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