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Key indicators

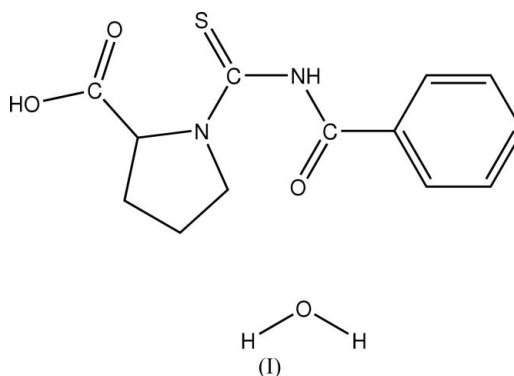
Single-crystal X-ray study
 $T = 298\text{ K}$
Mean $\sigma(\text{C}-\text{C}) = 0.008\text{ \AA}$
Disorder in main residue
 R factor = 0.037
 wR factor = 0.088
Data-to-parameter ratio = 13.2For details of how these key indicators were
automatically derived from the article, see
<http://journals.iucr.org/e>.(2S)-1-(Benzoylthiocarbamoyl)pyrrolidine-
2-carboxylic acid monohydrate

In the title compound, $\text{C}_{13}\text{H}_{14}\text{N}_2\text{O}_3\text{S}\cdot\text{H}_2\text{O}$, the benzoyl group lies out of the plane of the N_2CS thiourea unit, having a $\text{C}_{\text{carbonyl}}-\text{N}-\text{C}-\text{S}$ torsion angle of $119.4(4)^\circ$. The crystal structure is stabilized by $\text{N}-\text{H}\cdots\text{O}$, $\text{O}-\text{H}\cdots\text{O}$, $\text{C}-\text{H}\cdots\text{O}$ and $\text{C}-\text{H}\cdots\text{S}$ intermolecular hydrogen bonds, forming a two-dimensional network.

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Comment

In the benzoylthiourea derivatives of amino acids such as (2S)-2-(3-benzoylthioureido)-3-methylbutyric acid (Ngah *et al.*, 2005), and 2-(3-benzoylthioureido)propionic acid (Ngah *et al.*, 2006), the N atom of the amino acid attached to the thione carbon has an H atom bonded to it. However, in the title molecule, the pyrrolidine N atom has no attached H atom. The $\text{C7}-\text{N1}-\text{C8}-\text{S1}$ torsion angle of $119.4(4)^\circ$ shows that the benzoyl group lies significantly out of the thiourea plane. As in most benzoylthiourea compounds, the benzoyl group is approximately *trans* to the thiono S atom across the $\text{C7}-\text{N1}$ bond. The bonds length and angles are in normal ranges (Allen *et al.*, 1987).



The crystal structure is stabilized by intermolecular $\text{N}-\text{H}\cdots\text{O}$, $\text{O}-\text{H}\cdots\text{O}$, $\text{C}-\text{H}\cdots\text{O}$ and $\text{C}-\text{H}\cdots\text{S}$ hydrogen bonds, forming a two-dimensional network (Fig. 2, Table 1).

Experimental

An equimolar solution of benzoyl isothiocyanate (0.03 mol, 5.04 g) and proline (0.03 mol, 3.45 g) in acetone (200 ml) was refluxed for 1 h in a two-necked round-bottomed flask. The mixture was filtered into a beaker containing some ice. The resulting white precipitate was washed with distilled water followed by cold acetone, then dried in a vacuum desiccator (yield 75%, 6.26 g). Recrystallization from methanol yielded colourless crystals suitable for X-ray analysis (m.p. 430.1–430.6 K). The water of crystallization comes from wet solvent or atmospheric moisture.

Crystal data

$C_{13}H_{14}N_2O_3 \cdot H_2O$

$M_r = 296.34$

Orthorhombic, $P2_12_12_1$

$a = 9.05 (4) \text{ \AA}$

$b = 12.00 (6) \text{ \AA}$

$c = 13.33 (6) \text{ \AA}$

$V = 1448 (12) \text{ \AA}^3$

$Z = 4$

$D_x = 1.360 \text{ Mg m}^{-3}$

Mo $K\alpha$ radiation

$\mu = 0.24 \text{ mm}^{-1}$

$T = 298 (2) \text{ K}$

Block, colourless

$0.50 \times 0.30 \times 0.10 \text{ mm}$

Data collection

Bruker SMART APEX CCD area-detector diffractometer

ω scans

Absorption correction: multi-scan

(SADABS; Bruker, 2000)

$T_{\min} = 0.890, T_{\max} = 0.977$

6893 measured reflections

2527 independent reflections

2047 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.039$

$\theta_{\max} = 25.0^\circ$

Refinement

Refinement on F^2

$R[F^2 > 2\sigma(F^2)] = 0.037$

$wR(F^2) = 0.088$

$S = 0.99$

2527 reflections

191 parameters

H atoms treated by a mixture of independent and constrained refinement

$w = 1/[\sigma^2(F_o^2) + (0.0468P)^2]$

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

$(\Delta/\sigma)_{\max} < 0.001$

$\Delta\rho_{\max} = 0.17 \text{ e \AA}^{-3}$

$\Delta\rho_{\min} = -0.15 \text{ e \AA}^{-3}$

Absolute structure: Flack (1983),

634 Friedel pairs

Flack parameter: $-0.01 (10)$

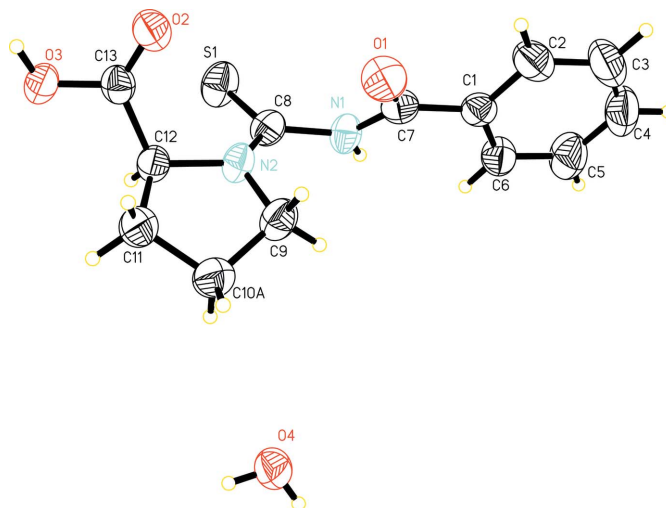


Figure 1

The molecular structure of the title compound, (I), shown with 50% probability displacement ellipsoids. The minor disorder component has been omitted.

Table 1

Hydrogen-bond geometry ($\text{\AA}, ^\circ$).

$D-H \cdots A$	$D-H$	$H \cdots A$	$D \cdots A$	$D-H \cdots A$
$N1-H1A \cdots O4^i$	0.86	2.31	3.113 (16)	156
$O3-H3 \cdots O4^{ii}$	0.82	1.97	2.754 (14)	160
$O4-H4A \cdots O2^{iii}$	0.82 (2)	2.22 (2)	2.945 (15)	148 (3)
$O4-H4B \cdots O1^{iii}$	0.82 (2)	1.91 (2)	2.728 (14)	177 (3)
$C11-H11A \cdots S1^{iv}$	0.97	2.87	3.754 (19)	152

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

H atoms of the water molecule were located in a difference map and refined freely. Other H atoms were positioned geometrically and allowed to ride on their parent atoms, with $C-H = 0.93-0.97 \text{ \AA}$, $O-H = 0.82 \text{ \AA}$ and $N-H = 0.86 \text{ \AA}$, and with $U_{\text{iso}}(H) = 1.2U_{\text{eq}}(C,N)$ for CH, CH_2 and NH, and $U_{\text{iso}}(H) = 1.5U_{\text{eq}}(O)$ for OH. Atom C10 and its attached H atoms are disordered over two sites, A and B, with site occupancy factors of 0.916 (7) and 0.084 (7), respectively.

Data collection: SMART (Bruker, 2000); cell refinement: SAINT (Bruker, 2000); 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, PARST (Nardelli, 1995) and PLATON (Spek, 2003).

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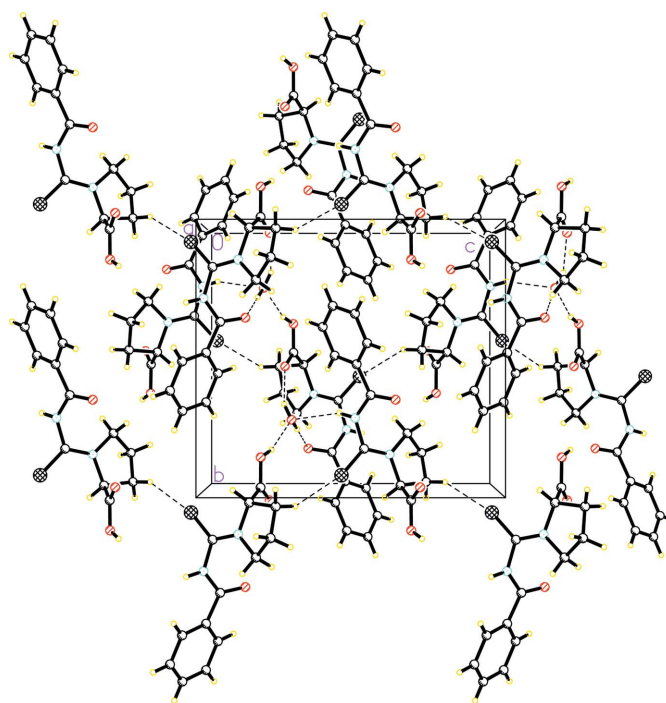


Figure 2

Packing diagram of (I), viewed down the a axis. The dashed lines denote hydrogen bonds. The minor disorder component has been omitted.

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