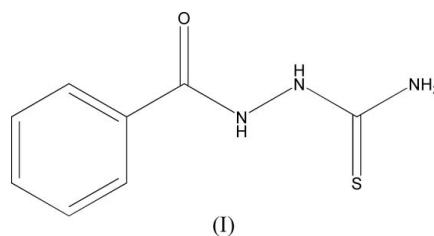


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541004, People's Republic of ChinaCorrespondence e-mail:  
bianhd@mailbox.gxnu.edu.cn**Key indicators**Single-crystal X-ray study  
 $T = 293$  K  
Mean  $\sigma(C-C) = 0.006$  Å  
 $R$  factor = 0.042  
 $wR$  factor = 0.124  
Data-to-parameter ratio = 12.8For details of how these key indicators were  
automatically derived from the article, see  
<http://journals.iucr.org/e>.***N*-Benzoylthiosemicarbazide**The title compound,  $C_8H_9N_3OS$ , crystallizes in a triclinic unit cell, with two crystallographically independent molecules in the asymmetric unit. It has a two-dimensional layer structure with intermolecular hydrogen bonding.

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**Comment***N*-Substituted thiosemicarbazides have been widely investigated because of the large number of potentially useful biological properties, particularly antitumour activity (Castiñeiras *et al.*, 2000). Therefore, we have synthesized the novel analogous compound *N*-benzoylthiosemicarbazide, (I), and determined its crystal structure.

The crystal structure of (I) is shown in Fig. 1. The hydrogen-bonding data for (I) are listed in Table 1. The title compound crystallizes in the triclinic space group  $P\bar{1}$ , with two crystallographically independent molecules in the asymmetric unit. There are two planes (plane 1: thiourea group; plane 2: benzoyl group) in each molecule. The angles between the two planes in the two molecules are  $56.9(3)$  and  $79.5(3)^\circ$ . In the two molecules, the angles at C7, C8, C15 and C16 are near  $120^\circ$ , indicating a trigonal planar arrangement around these C atoms. The angle between the two benzene rings is  $28.2(2)^\circ$ . The C8–S1 [ $1.721(2)$  Å] and C16–S2 [ $1.682(3)$  Å] bond lengths indicate character intermediate between single and double bonds, being shorter than a pure single bond (1.84 Å; Liu *et al.*, 1999).

In addition, neighbouring molecules are connected by hydrogen bonds, forming an infinite two-dimensional system (Fig. 2). Both molecules form intramolecular hydrogen bonds, *viz.* N3–H3A...N1 and N6–H6A...N4 in the two molecules. The S atoms interact with a total of five H atoms from five different molecules. Atom S1 interacts with the N1–H1, N5–H5 and N6–H6B groups, and S2 interacts with the N2–H2 and N3–H3B groups.

**Experimental**

Benzoyl chloride (39.5 mmol) was added to a solution of sodium benzoate (39.5 mmol) in chloroform (60 ml) at 278 K. The reaction mixture was then warmed slowly to 295 K and stirred for 2 h. Thio-

semicarbazide (32.9 mmol) was added to the mixture to give a white suspension, which was collected, rinsed with chloroform and diethyl ether, and then dried to give *N*-benzoylthiosemicarbazide. Colourless single crystals were obtained from absolute methanol.

Crystal data

$C_8H_9N_3OS$   $Z = 4$   
 $M_r = 195.24$   $D_x = 1.331 \text{ Mg m}^{-3}$   
 Triclinic,  $P\bar{1}$  Mo  $K\alpha$  radiation  
 Cell parameters from 905 reflections  
 $a = 8.237 (3) \text{ \AA}$   $\theta = 2.6\text{--}26.1^\circ$   
 $b = 8.438 (3) \text{ \AA}$   $\mu = 0.30 \text{ mm}^{-1}$   
 $c = 14.590 (6) \text{ \AA}$   $T = 293 (2) \text{ K}$   
 $\alpha = 83.554 (7)^\circ$  Block, colourless  
 $\beta = 75.631 (7)^\circ$   $0.20 \times 0.16 \times 0.14 \text{ mm}$   
 $\gamma = 84.998 (7)^\circ$   
 $V = 974.3 (7) \text{ \AA}^3$

Data collection

Bruker SMART CCD area-detector diffractometer 3408 independent reflections  
 $\varphi$  and  $\omega$  scans 2413 reflections with  $I > 2\sigma(I)$   
 Absorption correction: multi-scan (*SADABS*; Bruker, 1997)  $R_{int} = 0.020$   
 $T_{min} = 0.927, T_{max} = 0.959$   $\theta_{max} = 25.0^\circ$   
 5064 measured reflections  $h = -9 \rightarrow 9$   
 $k = -10 \rightarrow 8$   
 $l = -17 \rightarrow 11$

Refinement

Refinement on  $F^2$  H atoms treated by a mixture of independent and constrained refinement  
 $R[F^2 > 2\sigma(F^2)] = 0.042$   
 $wR(F^2) = 0.124$   
 $S = 1.09$   
 3408 reflections  $w = 1/[\sigma^2(F_o^2) + (0.066P)^2]$   
 267 parameters where  $P = (F_o^2 + 2F_c^2)/3$   
 $(\Delta/\sigma)_{max} = 0.001$   
 $\Delta\rho_{max} = 0.33 \text{ e \AA}^{-3}$   
 $\Delta\rho_{min} = -0.29 \text{ e \AA}^{-3}$

Table 1

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

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
$N6-H6B\cdots S1^i$	0.87 (1)	2.584 (12)	3.439 (3)	169 (3)
$N5-H5\cdots S1^{ii}$	0.85 (1)	2.650 (11)	3.495 (3)	172 (2)
$N4-H4\cdots S1^{iii}$	0.85 (1)	2.99 (3)	3.503 (3)	120 (3)
$N4-H4\cdots S2^{iv}$	0.85 (1)	2.90 (2)	3.584 (3)	138 (3)
$N3-H3B\cdots S2^v$	0.85 (1)	2.635 (12)	3.475 (3)	167 (3)
$N3-H3A\cdots O2^{vi}$	0.86 (1)	2.069 (13)	2.907 (3)	164 (3)
$N2-H2\cdots S2^{vii}$	0.86 (1)	2.431 (13)	3.268 (3)	165 (3)
$N1-H1\cdots S1^{viii}$	0.86 (1)	2.530 (13)	3.358 (3)	162 (2)

Symmetry codes: (i)  $1+x, 1+y, z-1$ ; (ii)  $1+x, y, z-1$ ; (iii)  $1-x, 1-y, 1-z$ ; (iv)  $2-x, 2-y, -z$ ; (v)  $x-1, y-1, 1+z$ ; (vi)  $2-x, 1-y, 1-z$ ; (vii)  $x-1, y, 1+z$ ; (viii)  $1-x, 1-y, 2-z$ .

The H atoms attached to N were located in a difference Fourier map and their coordinates and isotropic displacement parameters were refined [ $N-H = 0.85 (1)\text{--}0.87 (1) \text{ \AA}$ ]. All other H atoms were placed in idealized positions and were refined using a riding model [ $C-H = 0.93 \text{ \AA}$ ;  $U_{iso}(H) = 1.2U_{eq}(C)$ ].

Data collection: *SMART* (Bruker, 1998); cell refinement: *SAINT* (Bruker, 1998); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1990); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *SHELXTL* (Bruker, 1997); software used to prepare material for publication: *SHELXTL*.

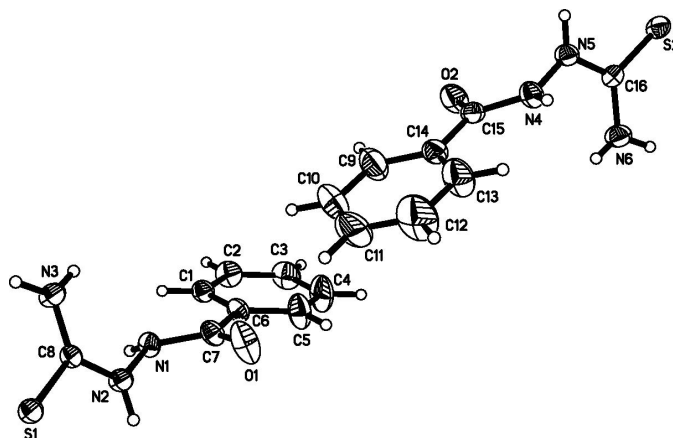


Figure 1

A view of the asymmetric unit of (I), showing the atom-numbering scheme. Displacement ellipsoids are drawn at the 50% level.

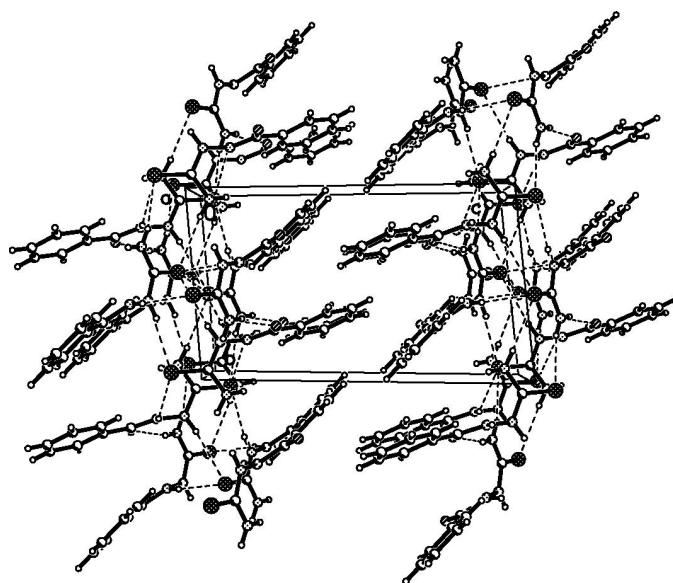


Figure 2

The two-dimensional network of hydrogen bonds (shown as dashed lines) in (I).

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