

Bohari M. Yamin\* and  
M. Sukeri M. YusofSchool of Chemical Sciences and Food  
Technology, Universiti Kebangsaan Malaysia,  
43600 Bangi, Selangor, MalaysiaCorrespondence e-mail:  
bohari@pkriscc.ukm.my

## Key indicators

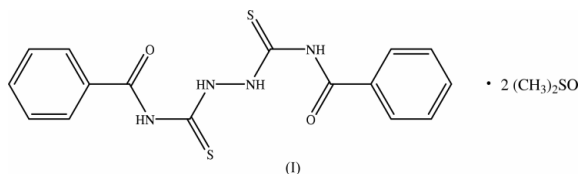
Single-crystal X-ray study  
 $T = 273\text{ K}$   
Mean  $\sigma(\text{C}-\text{C}) = 0.004\text{ \AA}$   
 $R$  factor = 0.055  
 $wR$  factor = 0.135  
Data-to-parameter ratio = 18.5For details of how these key indicators were  
automatically derived from the article, see  
<http://journals.iucr.org/e>.*N,N'*-Bis(benzamidothiocarbonyl)hydrazine  
dimethyl sulfoxide disolvate

In the title compound,  $\text{C}_{16}\text{H}_{14}\text{N}_4\text{O}_2\text{S}_2 \cdot 2\text{C}_2\text{H}_6\text{OS}$ , the molecule is discrete and centrosymmetric. The benzoylthiourea and benzoyl groups are *cis* and *trans*, respectively, to the thiourea S atom across the C–N bonds. In the crystal structure, the molecule is linked to the solvent DMSO molecules by N–H···O hydrogen bonds.

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## Comment

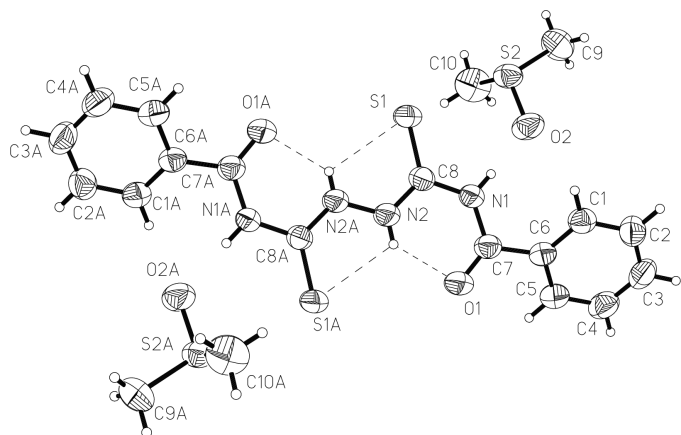
Benzoylthiourea derivatives can be synthesized from the reaction between benzoylisothiocyanate and amine compounds. Consequently, with a diamine, a bis(benzoylthiourea) derivative can be obtained. The title compound, (I), is a simple analogue of the derivative when hydrazine is used. The molecular structure and bond dimensions of (I) are in agreement with other benzoylthiourea derivatives, such as *N*-phenyl-*N'*-benzoylthiourea (Yamin & Yusof, 2003; Usman *et al.*, 2002; Kaminsky *et al.*, 2001). However, the molecule is centrosymmetric about the mid-point of the N2–N2<sup>i</sup> bond (Fig. 1 and Table 1; symmetry codes are defined in Table 1). The benzoylthiourea and benzoyl groups are *cis* and *trans* to atom S1 across the C–N bonds. The central bis-thiourea fragment, C6 to C6A, and the phenyl moieties are planar. The whole molecule looks like a wing with a dihedral angle between the central thiourea and each phenyl ring of 35.27 (9)°.



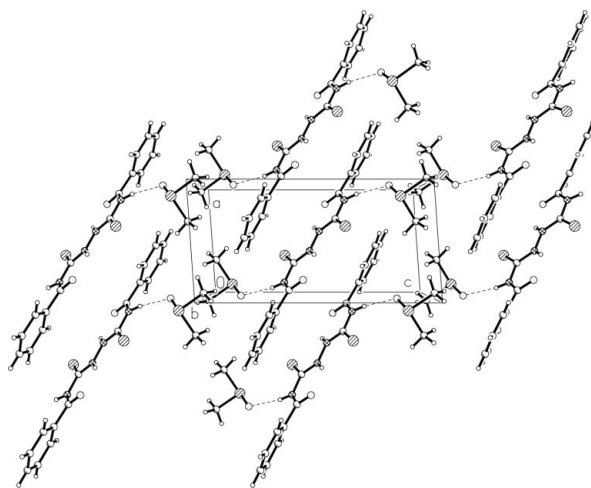
The presence of intramolecular hydrogen bonding, N2–H2A···O1 and N2–H2A···S1<sup>i</sup>, results in the formation of two pseudo-six-membered (N2/C8/N1/C7/O1/H2A) and five-membered rings (S1/C8/N2/N2A/H2A<sup>i</sup>) (Table 2). In the crystal structure, the molecule is linked to the solvent dimethyl sulfoxide (DMSO) molecules by N1–H1A···O2<sup>ii</sup> intermolecular hydrogen bonds (Fig. 2 and Table 2).

## Experimental

A solution of hydrazine (0.50 g, 9.9 mmol) in acetone (50 ml) was added dropwise to 50 ml of an acetone solution containing an equimolar amount of benzoylisothiocyanate in a two-neck round-bottomed flask. The solution was refluxed for about 2 h and then cooled in ice. The white precipitate was filtered off and washed with ethanol–distilled water, then dried in a vacuum (yield 80%). Recrystallization from DMSO yielded single crystals of (I) suitable for X-ray analysis.



**Figure 1**  
The molecular structure of the title compound, (I), shown with 50% probability displacement ellipsoids. Atoms labelled with a suffix 'A' are generated by the symmetry code  $-x, -y, -z$ .



**Figure 2**  
Packing diagram of (I), viewed down the  $b$  axis. The dashed lines denote  $N-H \cdots O$  hydrogen bonds.

#### Crystal data

$C_{16}H_{14}N_4O_2S_2 \cdot 2C_2H_6OS$   
 $M_r = 514.70$   
 Monoclinic,  $P2_1/n$   
 $a = 6.4127$  (11) Å  
 $b = 15.486$  (3) Å  
 $c = 12.805$  (2) Å  
 $\beta = 93.971$  (3)°  
 $V = 1268.6$  (4) Å<sup>3</sup>  
 $Z = 2$

#### Data collection

Bruker SMART APEX CCD area-detector diffractometer  
 $\omega$  scans  
 Absorption correction: multi-scan (SADABS; Sheldrick, 1996)  
 $T_{\min} = 0.804$ ,  $T_{\max} = 0.877$   
 7458 measured reflections

$D_x = 1.347$  Mg m<sup>-3</sup>  
 Mo  $K\alpha$  radiation  
 Cell parameters from 1464 reflections  
 $\theta = 2.1$ – $27.5^\circ$   
 $\mu = 0.41$  mm<sup>-1</sup>  
 $T = 273$  (2) K  
 Block, colourless  
 $0.56 \times 0.34 \times 0.33$  mm

2862 independent reflections  
 2008 reflections with  $I > 2\sigma(I)$   
 $R_{\text{int}} = 0.034$   
 $\theta_{\text{max}} = 27.5^\circ$   
 $h = -6 \rightarrow 8$   
 $k = -20 \rightarrow 18$   
 $l = -16 \rightarrow 16$

#### Refinement

Refinement on  $F^2$   
 $R[F^2 > 2\sigma(F^2)] = 0.055$   
 $wR(F^2) = 0.135$   
 $S = 1.03$   
 2862 reflections  
 155 parameters  
 H atoms treated by a mixture of independent and constrained refinement

$$w = 1/[\sigma^2(F_o^2) + (0.0599P)^2 + 0.1638P]$$

$$\text{where } P = (F_o^2 + 2F_c^2)/3$$

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

$$\Delta\rho_{\text{max}} = 0.39 \text{ e \AA}^{-3}$$

$$\Delta\rho_{\text{min}} = -0.18 \text{ e \AA}^{-3}$$

**Table 1**

Selected geometric parameters (Å, °).

C8—N2	1.323 (3)	C9—S2	1.780 (3)
C8—N1	1.386 (3)	C10—S2	1.764 (3)
C8—S1	1.664 (2)	N2—N2 <sup>i</sup>	1.372 (4)
C7—O1	1.224 (3)	O2—S2	1.5026 (19)
C7—N1	1.380 (3)		
N2—C8—N1	116.0 (2)	N1—C8—S1	120.94 (18)
N2—C8—S1	123.1 (2)	O1—C7—N1	122.5 (2)

Symmetry codes: (i)  $-x, -y, -z$ .

**Table 2**

Hydrogen-bonding geometry (Å, °).

$D-H \cdots A$	$D-H$	$H \cdots A$	$D \cdots A$	$D-H \cdots A$
N2—H2A <sup>i</sup> ···O1	0.78 (3)	2.01 (3)	2.593 (3)	132 (2)
N2—H2A <sup>i</sup> ···S1 <sup>i</sup>	0.78 (3)	2.53 (3)	2.920 (2)	113 (2)
N1—H1A <sup>ii</sup> ···O2 <sup>ii</sup>	0.84 (3)	2.01 (3)	2.843 (3)	169 (3)

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

H atoms bonded to N atoms were located from difference maps and refined isotropically. H atoms bonded to C atoms were fixed geometrically at their ideal positions and allowed to ride on the parent atoms, with  $C-H = 0.97$  Å.

Data collection: SMART (Siemens, 1996); cell refinement: SAINT (Siemens, 1996); data reduction: SAINT; program(s) used to solve structure: SHELXTL (Sheldrick, 1997); program(s) used to refine structure: SHELXTL; molecular graphics: SHELXTL; software used to prepare material for publication: SHELXTL, PARST (Nardelli, 1995) and PLATON (Spek, 1990).

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