

*N'*-Benzoyl-*N*-*p*-bromophenylthioureaBohari M. Yamin\* and  
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## Key indicators

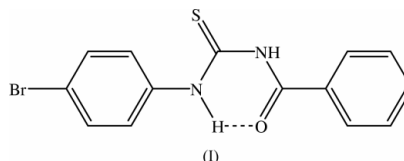
Single-crystal X-ray study  
 $T = 273\text{ K}$   
Mean  $\sigma(\text{C}-\text{C}) = 0.004\text{ \AA}$   
 $R$  factor = 0.033  
 $wR$  factor = 0.091  
Data-to-parameter ratio = 17.9For details of how these key indicators were  
automatically derived from the article, see  
<http://journals.iucr.org/e>.

In the title compound,  $\text{C}_{14}\text{H}_{11}\text{BrN}_2\text{OS}$ , the bromophenyl and benzoyl groups lie *cis* and *trans*, respectively, to the S atom across the thiourea C–N bonds. Owing to the presence of the Br atom in the *para* position, the dihedral angle between the bromophenyl group and the central carbonylthiourea plane is increased to  $20.40(11)^\circ$ , in comparison with  $7.52(9)^\circ$  in its unsubstituted phenyl analogue. The molecules are linked by intermolecular contacts  $\text{N}-\text{H}\cdots\text{S}$  and  $\text{C}-\text{H}\cdots\text{O}$  to form linear chains parallel to the *a* axis.

Received 21 January 2003  
Accepted 11 February 2003  
Online 21 February 2003

## Comment

Benzoylthiourea derivatives adopt a *cis*–*trans* conformation with respect to the position of the substituent and benzoyl groups relative to the S atom across the C–N bonds (Shanmuga Sundara Raj *et al.*, 1999; Usman *et al.*, 2002; Kaminsky *et al.*, 2002). The structure and bond dimensions of the title compound, (I), are in agreement with those found in aryl-benzoylthioureas, including *N*-phenyl-*N'*-benzoylthiourea, (II) (Yamin & Yusof, 2003). However, the presence of Br1 at the *para* position causes the dihedral angle between the bromophenyl plane (C9–C14/Br1) and the central thiourea fragment (S1/C8/N1/N2/C7/O1) to increase from  $7.52(9)^\circ$  in (II) to  $20.40(11)^\circ$  (Table 1). The phenyl (C1–C6) plane and the bromophenyl plane make an angle of  $38.61(11)^\circ$ , compared to  $33.3(1)^\circ$  in (II) (Yamin & Yusof, 2003). The dihedral angle between the phenyl and the central thiourea fragment is  $29.93(11)^\circ$ . As in (II), there are two intramolecular hydrogen bonds,  $\text{C14}-\text{H14A}\cdots\text{S1}$  and  $\text{N2}-\text{H2A}\cdots\text{O1}$ , maintaining the presence of a pseudo-six-membered ring (N1–C8–N2–H2A $\cdots$ O1–C7) (Table 2). However, the molecules are packed through weak  $\text{N1}-\text{H1B}\cdots\text{S1}^i$  and  $\text{C11}-\text{H11}\cdots\text{O1}^{ii}$  [symmetry codes: (i)  $1-x, -y, 2-z$ ; (ii)  $-x, -y, 2-z$ ] intermolecular contacts to form a linear chain parallel to the *a* axis (Fig. 2).



## Experimental

A solution of *para*-bromoaniline (0.50 g, 2.9 mmol) in acetone (50 ml) was added dropwise to 50 ml of an acetone solution containing an equimolar amount of benzoyl thiocyanate in a two-neck round-bottomed flask. The solution was refluxed for about 1 h and then cooled in ice. The white precipitate which formed was filtered off and washed with ethanol–distilled water, then dried in a vacuum (yield 79%). Recrystallization from DMSO yielded single

crystals suitable for X-ray analysis.

### Crystal data

$C_{14}H_{11}BrN_2OS$   
 $M_r = 335.22$   
 Monoclinic,  $P2_1/c$   
 $a = 13.846$  (3) Å  
 $b = 5.9486$  (14) Å  
 $c = 16.972$  (4) Å  
 $\beta = 103.311$  (4)°  
 $V = 1360.3$  (6) Å<sup>3</sup>  
 $Z = 4$

$D_x = 1.637$  Mg m<sup>-3</sup>  
 Mo  $K\alpha$  radiation  
 Cell parameters from 2075 reflections  
 $\theta = 1.5$ – $27.5$ °  
 $\mu = 3.16$  mm<sup>-1</sup>  
 $T = 273$  (2) K  
 Block, colourless  
 $0.33 \times 0.30 \times 0.22$  mm

### Data collection

Bruker SMART APEX CCD area-detector  
 $\omega$  scans  
 Absorption correction: multi-scan (SADABS; Sheldrick, 1996)  
 $T_{\min} = 0.421$ ,  $T_{\max} = 0.542$   
 7554 measured reflections

3081 independent reflections  
 2284 reflections with  $I > 2\sigma(I)$   
 $R_{\text{int}} = 0.026$   
 $\theta_{\text{max}} = 27.5$ °  
 $h = -12 \rightarrow 17$   
 $k = -7 \rightarrow 7$   
 $l = -21 \rightarrow 22$

### Refinement

Refinement on  $F^2$   
 $R[F^2 > 2\sigma(F^2)] = 0.033$   
 $wR(F^2) = 0.091$   
 $S = 0.91$   
 3081 reflections  
 172 parameters  
 H-atom parameters constrained

$w = 1/[\sigma^2(F_o^2) + (0.0511P)^2 + 0.2498P]$   
 where  $P = (F_o^2 + 2F_c^2)/3$   
 $(\Delta/\sigma)_{\text{max}} < 0.001$   
 $\Delta\rho_{\text{max}} = 0.41$  e Å<sup>-3</sup>  
 $\Delta\rho_{\text{min}} = -0.48$  e Å<sup>-3</sup>

**Table 1**

Selected geometric parameters (Å, °).

Br1–C12	1.900 (2)	O1–C7	1.220 (3)
S1–C8	1.661 (3)	C7–N1	1.372 (3)
N2–C8	1.325 (3)	C8–N1	1.394 (3)
N2–C9	1.418 (3)		
C8–N2–C9	132.1 (2)	N2–C8–N1	114.3 (2)
C13–C12–Br1	119.47 (19)	N2–C8–S1	127.57 (19)
O1–C7–N1	122.1 (2)	N1–C8–S1	118.15 (18)

**Table 2**

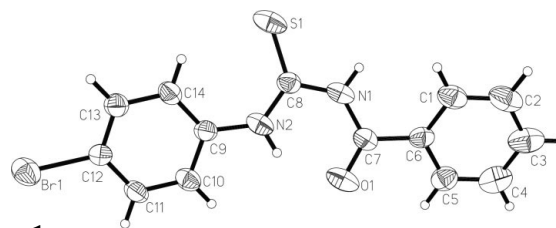
Hydrogen-bonding geometry (Å, °).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
N2–H2A <sup>i</sup> ⋯O1	0.86	1.87	2.606 (3)	143
C14–H14A <sup>i</sup> ⋯S1	0.93	2.64	3.246 (3)	123
N1–H1B <sup>i</sup> ⋯S1 <sup>i</sup>	0.86	2.72	3.534 (2)	158
C11–H11A <sup>i</sup> ⋯O1 <sup>ii</sup>	0.93	2.51	3.421 (3)	165

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

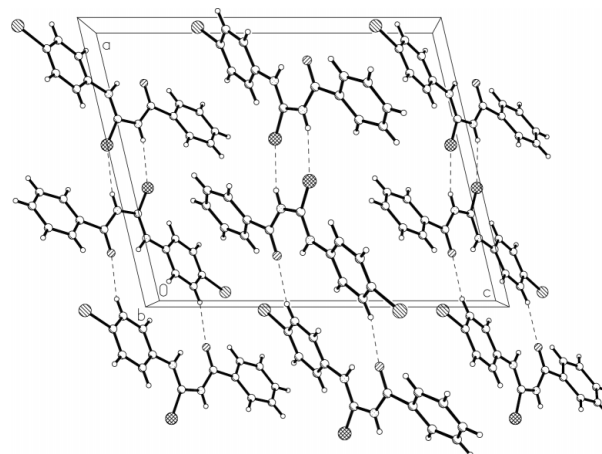
After their location in a difference map, all H atoms were included in the refinement in geometrically determined positions and made to ride on the parent C or N atoms, with C–H = 0.97 Å and N–H = 0.89 Å.

Data collection: *SMART* (Siemens, 1996); cell refinement: *SAINT* (Siemens, 1996); data reduction: *SAINT*; program(s) used to solve



**Figure 1**

The molecular structure of the title compound, shown with 50% probability displacement ellipsoids.



**Figure 2**

Packing diagram of (I), viewed down the  $b$  axis. Dashed lines indicate N–H⋯S and C–H⋯O hydrogen bonds.

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).

The authors thank the Malaysian Government and Universiti Kebangsaan Malaysia for the research grants IRPA No. 09-02-02-0163.

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