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#### **Key indicators**

Single-crystal X-ray study T = 173 K Mean  $\sigma$ (C–C) = 0.002 Å R factor = 0.037 wR factor = 0.106 Data-to-parameter ratio = 22.4

For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e.

# 1-(2-Chlorobenzoyl)-3-phenylthiourea

The phenyl and benzoyl groups in the title molecule,  $C_{14}H_{11}CIN_2OS$ , are *cis* and *trans*, respectively, with respect to the C=S bond. The molecular conformation is stabilized by an N-H···O hydrogen bond and the crystal packing is characterized by N-H···O and N-H···S hydrogen bonds.

## Comment

The background to this study has been set out in the preceding paper (Rauf *et al.*, 2006).



The molecular structure of (I) is shown in Fig. 1. Bond lengths and angles can be regarded as normal (Cambridge Structural Database, Version 5.27 plus one update; *MOGUL* Version 1.1; Allen, 2002). The central carbonylthiourea group is almost planar (r.m.s. deviation for O1/C1/N1/C2/S1/N2 is 0.054 Å). The chlorophenyl and phenyl rings form dihedral angles of 74.31 (3) and 40.22 (4)°, respectively, with the central carbonylthiourea plane. The dihedral angle between the two aromatic rings is 34.92 (5)°. An intramolecular N-H···O hydrogen bond gives rise to a six-membered ring. The crystal packing is stabilized by N-H···O and N-H···S hydrogen bonds (Table 2).

# **Experimental**

A solution of 2-chlorobenzoyl chloride (1.750 g, 10 mmol) in acetone (50 ml) was added dropwise to a suspension of KSCN (1.00 g, 10 mmol) in acetone (30 ml). The reaction mixture was heated under reflux for 45 min, and then cooled to room temperature. Afterwards a solution of aniline (0.93 g, 10 mmol) in acetone (15 ml) was added



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Received 22 February 2006 Accepted 28 February 2006 and the resulting mixture was stirred for 3 h. The reaction mixture was then poured into crushed ice and stirred well. The solid product was separated and washed with deionized water and purified by recrystallization from toluene to give crystals of the title compound, in an overall yield of 85%.

 $D_r = 1.410 \text{ Mg m}^{-3}$ 

Cell parameters from 24244

Mo  $K\alpha$  radiation

reflections

 $\theta=2.5{-}30.4^\circ$ 

 $\mu = 0.42~\mathrm{mm}^{-1}$ 

T = 173 (2) K

Plate, colourless

 $0.37 \times 0.35 \times 0.17 \text{ mm}$ 

#### Crystal data

 $C_{14}H_{11}ClN_2OS$   $M_r = 290.76$ Monoclinic, C2/c a = 20.5099 (13) Å b = 7.3281 (3) Å c = 18.5344 (12) Å  $\beta = 100.437$  (5)° V = 2739.6 (3) Å<sup>3</sup> Z = 8

## Data collection

Stoe IPDS-II two-circle	4055 independent reflections
diffractometer	3793 reflections with $I > 2\sigma(I)$
$\omega$ scans	$R_{\rm int} = 0.059$
Absorption correction: multi-scan	$\theta_{\rm max} = 30.3^{\circ}$
(MULABS; Spek, 2003; Blessing,	$h = -28 \rightarrow 28$
1995)	$k = -10 \rightarrow 10$
$T_{\min} = 0.859, \ T_{\max} = 0.932$	$l = -22 \rightarrow 26$
24244 measured reflections	

#### Refinement

Refinement on $F^2$	$w = 1/[\sigma^2(F_0^2) + (0.0631P)^2]$
$R[F^2 > 2\sigma(F^2)] = 0.037$	+ 1.3534P]
$wR(F^2) = 0.106$	where $P = (F_0^2 + 2F_c^2)/3$
S = 1.07	$(\Delta/\sigma)_{\rm max} = 0.001$
4055 reflections	$\Delta \rho_{\rm max} = 0.40 \ {\rm e} \ {\rm \AA}^{-3}$
181 parameters	$\Delta \rho_{\rm min} = -0.50 \text{ e } \text{\AA}^{-3}$
H atoms treated by a mixture of	Extinction correction: SHELXL97
independent and constrained	Extinction coefficient: 0.0112 (10)
refinement	

#### Table 1

Selected bond lengths (Å).

Cl1-C12	1.7419 (13)	C1-N1	1.3725 (13
S1-C2	1.6758 (11)	C2-N2	1.3391 (13
C1 - O1	1 2269 (13)		

# Table 2

Hydrogen-bond geometry (Å, °).

$D - H \cdots A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdots A$
$N1-H1\cdots S1^{i}$ $N2-H2\cdots O1$ $N2-H2\cdots O1^{ii}$	0.903 (18)	2.492 (18)	3.3743 (9)	165.6 (16)
	0.89 (2)	1.95 (2)	2.6888 (12)	138.6 (17)
	0.89 (2)	2.57 (2)	3.2917 (13)	138.0 (15)

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

H atoms were located in a difference electron-density map, but those bonded to C atoms were refined with fixed individual displacement parameters  $[U_{iso}(H) = 1.2U_{eq}(C)]$  using a riding model, with C-H = 0.95 Å. The H atoms bonded to nitrogen were refined freely.

Data collection: X-AREA (Stoe & Cie, 2001); cell refinement: X-AREA; data reduction: X-AREA; program(s) used to solve structure: SHELXS97 (Sheldrick, 1990); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: XP in SHELXTL-Plus (Sheldrick, 1991); software used to prepare material for publication: SHELXL97 and PLATON (Spek, 2003).

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