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Key indicators

Single-crystal X-ray study T = 120 K Mean σ (C–C) = 0.004 Å R factor = 0.057 wR factor = 0.120 Data-to-parameter ratio = 16.7

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

1-(3-Chlorobenzoyl)-3-[3-(trifluoromethyl)phenyl]thiourea

In the crystal structure of the title compound, $C_{15}H_{10}ClF_3N_2OS$, the dihedral angle between the two aromatic rings is 32.68 (8)°. The crystal packing shows intermolecular N-H···O and C-H···S hydrogen bonds which link molecules into sheets which are stacked along [001].

Comment

N-Substituted and N,N'-disubstituted thiourea derivatives are very useful building blocks for the synthesis of a wide range of aliphatic macromolecular and heterocyclic compounds. Thus, benzothiazoles have been prepared from arylthioureas in the presence of bromine (Patil & Chedekel, 1984), and the condensation of thiourea with α -halocarbonyl compounds forms 2-aminothiazoles (Baily et al., 1996). 2-Methyl-aminothiazolines have been synthesized by cyclization of N-(2hydroxyethyl)-N'-methylthioureas (Namgun et al., 2001). Thioureas are efficient guanylating agents (Maryanoff et al., 1986). N.N-Dialkyl-N-aroylthioureas have been efficiently used for the extraction of nickel, palladium and platinum metals (Koch, 2001). Aliphatic and acylthioureas are well known for their fungicidal, antiviral, pesticidal and plantgrowth regulating activities (Upadlgaya & Srivastava, 1982; Wegner et al., 1986). Symmetrical and unsymmetrical thioureas have shown antifungal activity against the plant pathogens Pyricularia oryzae and Drechslera orvzae (Krishnamurthy et al., 1999). We became interested in the synthesis of these thioureas as intermediates in the synthesis of novel guanidines and heterocyclic compounds for the systematic study of bioactivity and complexation behaviour and we present here the crystal structure of the title compound, (I).



Compound I (Fig. 1) shows the typical thiourea C=S and C=O double bonds, as well as shortened C-N bonds (Table 1). The thiocarbonyl and carbonyl groups are almost coplanar, as reflected by the torsion angles $C8-N2-C9-O1 = 10.6 (5)^{\circ}$ and $C9-N2-C8-N1 = -18.1 (5)^{\circ}$. This is asso-

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Figure 1

The structure of (I), showing the atom-numbering scheme. Displacement ellipsoids are drawn at the 50% probability level.



Figure 2

The crystal packing of (I), viewed along [010], with hydrogen bonds indicated as dashed lines. H atoms not involved in hydrogen bonding have been omitted.

ciated with the expected typical thiourea intramolecular N1– $H1A\cdots O1$ hydrogen bond (Table 2). The dihedral angle formed by the two benzene ring planes, C1–C6 and C10–C15, is 32.68 (8)°. Other geometric parameters present no unusual features (Khawar Rauf, Badshah & Bolte, 2006; Khawar Rauf, Badshah & Flörke, 2006).

The crystal packing of (I) shows intermolecular $N-H\cdots O$ and $C-H\cdots S$ hydrogen bonds (Table 2, Fig. 2) which link the molecules into sheets stacked along [001]. The Cl atoms and N2-H group are not involved in any hydrogen bonds.

Experimental

Freshly prepared 3-chlorobenzoyl chloride (1.75 g, 10 m mol) was added to a suspension of KSCN ((1.00 g, 10 mmol) in acetone (30 ml). The reaction mixture was stirred for 15 min. Neat 3-(trifluoromethyl)aniline (1.6 g, 10 mmol) was then added and the resulting mixture was stirred for 1 h. The reaction mixture was then poured into acidified water (500 ml) and stirred well. The solid product was separated, washed with deionized water and purified by recrystallization from methanol–dichloromethane (1:1) to give fine crystals of (I), with an overall yield of 85%. The full spectroscopic and physical characterization of the compound will be reported elsewhere.

Crystal data

С

M M

a b c

β

v

$_{15}H_{10}ClF_3N_2OS$	Z = 4
$T_r = 358.76$	$D_x = 1.634 \text{ Mg m}^{-3}$
onoclinic, $P2_1/c$	Mo $K\alpha$ radiation
= 15.249 (4) Å	$\mu = 0.44 \text{ mm}^{-1}$
= 6.5459 (16) Å	T = 120 (2) K
= 14.643 (4) Å	Plate, colourless
= 94.066 (5)°	$0.48 \times 0.22 \times 0.03$
= 1457.9 (7) Å ³	

Data collection

Bruker SMART APEX CCD areadetector diffractometer φ and ω scans Absorption correction: multi-scan (*SADABS*; Bruker, 2002) $T_{\min} = 0.816, T_{\max} = 0.987$

Refinement

Refinement on F^2 $R[F^2 > 2\sigma(F^2)] = 0.057$ $wR(F^2) = 0.121$ S = 0.893468 reflections 208 parameters $0.48 \times 0.22 \times 0.03 \text{ mm}$

9777 measured reflections 3468 independent reflections 1968 reflections with $I > 2\sigma(I)$ $R_{int} = 0.094$ $\theta_{max} = 27.9^{\circ}$

H-atom parameters constrained
$w = 1/[\sigma^2 (F_o^2) + (0.0464P)^2]$
where $P = (F_0^2 + 2F_c^2)/3$
$(\Delta/\sigma)_{\rm max} < 0.001$
$\Delta \rho_{\rm max} = 0.94 \text{ e } \text{\AA}^{-3}$
$\Delta \rho_{\rm min} = -0.57 \text{ e } \text{\AA}^{-3}$

Table 1

Selected geometric parameters (Å, °).

Cl1-C12	1.736 (3)	N1-C1	1.438 (4)
S1-C8	1.667 (3)	N2-C9	1.375 (4)
O1-C9	1.217 (4)	N2-C8	1.392 (4)
N1-C8	1.326 (4)	C9-C10	1.488 (4)
C8-N1-C1	121.3 (3)	N2-C8-S1	117.5 (2)
C9-N2-C8	128.1 (3)	O1-C9-N2	122.3 (3)
N1-C8-N2	116.8 (3)	O1-C9-C10	123.1 (3)
N1-C8-S1	125.7 (2)	N2-C9-C10	114.5 (3)

Table 2			
Hydrogen-bond	geometry	(Å,	°).

$D - H \cdots A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdot \cdot \cdot A$
N1−H1A···O1	0.88	2.05	2.683 (3)	128
$N1 - H1A \cdots O1^{i}$	0.88	2.53	3.199 (3)	133
$C2-H2A\cdots S1^{ii}$	0.95	2.93	3.599 (3)	128
$C15-H15A\cdots S1^{iii}$	0.95	2.84	3.666 (3)	146

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

H atoms were located in a difference Fourier map and refined in idealized positions riding on their parent C and N atoms, with C–H = 0.95 Å and N–H = 0.88 Å, and with isotropic displacement parameters $U_{\rm iso}({\rm H}) = 1.2U_{\rm eq}({\rm C,N})$. The anisotropic displacement parameters of the trifluoromethyl group indicate some degree of disorder (rotation along the C5–C7 axis) but this could not be resolved.

Data collection: *SMART* (Bruker, 2002); cell refinement: *SAINT* (Bruker, 2002); data reduction: *SAINT*; program(s) used to solve structure: *SHELXTL* (Bruker, 2002); program(s) used to refine structure: *SHELXTL*; molecular graphics: *SHELXTL*; software used to prepare material for publication: *SHELXTL*.

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