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#### **Key indicators**

Single-crystal X-ray study T = 293 K Mean  $\sigma$ (C–C) = 0.003 Å R factor = 0.042 wR factor = 0.105 Data-to-parameter ratio = 14.3

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

# N-(3,4-Dichlorophenyl)-N'-(3-nitrobenzoyl)thiourea

In the title molecule,  $C_{14}H_9N_3O_3Cl_2S$ , the dihedral angles between the mean plane of the central carbonylthiourea group, N<sub>2</sub>CSCO, and the mean planes of the 3-nitrobenzoyl and 3,4-dichlorophenyl groups are 22.82 (9) and 69.75 (11)°, respectively. The crystal packing is stabilized by weak intermolecular C-H···O hydrogen bonds, forming a onedimensional chain along the *b* axis.

#### Comment

The title compound, (I), is analogous to N-(4-chloro-3-nitrophenyl)-N'-(3-nitrobenzoyl)thiourea, (II) (Yusof et al., 2006), except that the bulky nitro group in the disubstituted benzene ring is replaced by a Cl atom (Fig.1). The molecule maintains its trans-cis configuration with respect to the position of the 3nitrobenzoyl and 3,4-dichlorophenyl groups relative to the thiono S1 atom across the C8-N2 and C8-N3 bonds, respectively. The bond lengths and angles are in normal ranges (Allen et al., 1987) and are comparable to those in (II). The central carbonylthiourea (S1/C8/N2/N3/C7/O3), 3-nitrophenyl (C1-C6/N1/O1/O2) and 3,4-dichlorophenyl (C9-C14/Cl1/Cl2) fragments are all planar, with a maximum deviation of 0.107 (3)Å for atom O2. The mean plane of the central carbonylthiourea group, N<sub>2</sub>CSCO, makes dihedral angles of 22.82 (9) and 69.75  $(11)^{\circ}$ , respectively, with the mean planes of the 3-nitrobenzoyl and 3,4-dichlorophenyl groups. The dihedral angle between the two aromatic rings is  $48.25 (10)^{\circ}$ .



There are two intramolecular hydrogen bonds, N3– H3...O3 and C14–H14...S1 (Table 2), forming two pseudosix-membered rings, *viz.* N3–H3...O3–C7–N2–C8–N3 and C14–H14...S1–C8–N3–C9–C14. In the crystal structure, the molecules are linked by intermolecular interactions, C1–H1A...O2<sup>i</sup> (symmetry code as in Table 2), forming a one-dimensional chain along the *b* axis (Fig. 2).

## Experimental

An equimolar amount of 3,4-dichloroaniline (1.78 g, 11 mmol) in 20 ml acetone was added dropwise to a stirred acetone solution (75 ml) containing 3-nitrobenzoyl chloride (2.0 g, 11 mmol) and

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## Figure 1

The molecular structure of the title compound, (I), shown with 50% probability displacement ellipsoids. Hydrogen bonds are shown as dashed lines.



#### Figure 2

Packing diagram for compound (I), viewed down the *a* axis. Dashed lines denote  $C-H \cdots O$  hydrogen bonds.

ammonium thiocyanate (0.82 g, 11 mmol). The solution mixture was refluxed for 1 h. The resulting solution was poured into a beaker containing some ice blocks. The light-yellow precipitate was filtered off and washed with distilled water and cold ethanol before being dried under vacuum. Good quality crystals were obtained by recrystallization from methanol (yield 67%, 2.73 g; m.p. 489.3–491.7 K).

#### Crystal data

| $C_{14}H_9Cl_2N_3O_3S$          | V = 762.0 (3) Å <sup>3</sup>              |
|---------------------------------|---|
| $M_r = 370.20$                  | Z = 2                                     |
| Triclinic, P1                   | $D_x = 1.613 \text{ Mg m}^{-3}$           |
| a = 7.1648 (15)  Å              | Mo $K\alpha$ radiation                    |
| $b = 8.1522 (17) \text{\AA}$    | $\mu = 0.58 \text{ mm}^{-1}$              |
| c = 13.866 (3) Å                | T = 293 (2) K                             |
| $\alpha = 73.990 \ (3)^{\circ}$ | Slab, light yellow                        |
| $\beta = 81.912 \ (3)^{\circ}$  | $0.50 \times 0.22 \times 0.08 \text{ mm}$ |
| $\nu = 79.552 \ (4)^{\circ}$    |   |

#### Data collection

| Bruker SMART APEX CCD area-            |  |
|--|--|
| detector diffractometer                |  |
| w scans                                |  |
| Absorption correction: multi-scan      |  |
| (SADABS; Bruker, 2000)                 |  |
| $T_{\min} = 0.760, \ T_{\max} = 0.955$ |  |

#### Refinement

| Refinement on $F^2$             | $w = 1/[\sigma^2(F_0^2) + (0.0508P)^2]$                    |
|---------------------------------|--|
| $R[F^2 > 2\sigma(F^2)] = 0.042$ | + 0.2184P]   |
| $wR(F^2) = 0.105$               | where $P = (F_0^2 + 2F_c^2)/3$                             |
| S = 1.04                        | $(\Delta/\sigma)_{\rm max} < 0.001$                        |
| 2964 reflections                | $\Delta \rho_{\rm max} = 0.30 \text{ e} \text{ \AA}^{-3}$  |
| 208 parameters                  | $\Delta \rho_{\rm min} = -0.26 \text{ e } \text{\AA}^{-3}$ |
| H-atom parameters constrained   |  |

7867 measured reflections 2964 independent reflections 2400 reflections with  $I > 2\sigma(I)$ 

 $R_{\rm int} = 0.023$  $\theta_{\rm max} = 26.0^{\circ}$ 

#### Table 1

Selected geometric parameters (Å, °).

| Cl1-Cl2  | 1.730 (2) | S1-C8    | 1.648 (2)   |
|----------|-----------|----------|-------------|
| Cl2-Cl3  | 1.735 (2) | N2-C8    | 1.397 (3)   |
| N2-C7-C6 | 116.1 (2) | N3-C8-S1 | 126.61 (16) |
| N3-C8-N2 | 114.6 (2) | N2-C8-S1 | 118.80 (17) |

## Table 2

D N C

Hydrogen-bond geometry (Å, °).

| $-\mathrm{H}\cdots A$  | D-H                  | $H \cdot \cdot \cdot A$ | $D \cdots A$                        | $D - \mathbf{H} \cdots A$ |
|--|----------------------|-------------------------|-------------------------------------|---------------------------|
| $ \begin{array}{l} 3-H3\cdotsO3\\ 14-H14\cdotsS1\\ 1-H1A\cdotsO2^{i} \end{array} $ | 0.86<br>0.93<br>0.93 | 1.94<br>2.83<br>2.42    | 2.648 (3)<br>3.225 (2)<br>3.322 (4) | 138<br>107<br>165         |
|  |                      |                         |                                     |                           |

Symmetry code: (i) x, y - 1, z.

After their location in a difference map, all H atoms were positioned geometrically, with N–H = 0.86 Å and C–H = 0.93–0.96 Å, and constrained to ride on their parent atoms, with  $U_{iso}(H)$ =  $xU_{eq}(C,N)$ , where x = 1.5 for methyl H and 1.2 for all other H atoms.

Data collection: *SMART* (Bruker, 2000); cell refinement: *SAINT* (Bruker, 2000); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997*a*); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997*a*); molecular graphics: *SHELXTL* (Sheldrick, 1997*b*); software used to prepare material for publication: *SHELXTL*, *PARST* (Nardelli, 1995) and *PLATON* (Spek, 2003).

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