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#### **Key indicators**

Single-crystal X-ray study T = 293 K Mean  $\sigma$ (C–C) = 0.009 Å R factor = 0.057 wR factor = 0.145 Data-to-parameter ratio = 16.0

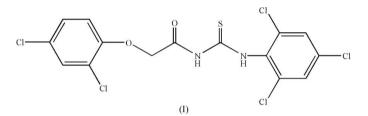
For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e.

# *N*-(2,4-Dichlorophenoxyacetyl)-*N*'-(2,4,6-trichlorophenyl)thiourea

In the title molecule,  $C_{15}H_9Cl_5N_2O_2S$ , all bond lengths and angles are normal. Weak intermolecular  $C-H\cdots S$  hydrogen bonds link the molecules into spiral chains running along the *c* axis. The crystal packing exhibits a short  $Cl\cdots Cl$  distance of 3.364 (3) Å, which suggests the existence of intermolecular  $Cl\cdots Cl$  interactions.

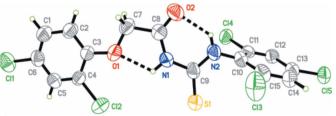
#### Comment

We have previously reported the structure of N-(2,4-dichlorophenoxyacetyl)-N'-(4-nitrophenyl)thiourea, (II) (Li, Yang, Xie & Luo, 2006). In continuation of our study of aroylthiourea compounds, we report here the crystal structure of the title compound, (I).



In (I) (Fig. 1), all bond lengths and angles agree with those observed in related compounds (Li, Yang, Xie & Luo, 2006; Li, Yang & Xie, 2006). The mean planes C1–C6/C11/Cl2 (*A*), C7–C9/O1/O2/N1/N2/S1 (*B*) and C10–C15/Cl3–Cl5 (*C*) make the following dihedral angles: A/B = 12.15 (16)° [1.10 (10)° in (II)], A/C = 81.58 (14)° [9.96 (13)° in (II)] and B/C = 80.70 (13)° [8.86 (12)° in (II)]. The intramolecular N–H···O hydrogen bonds (Table 1) influence the molecular conformation.

The weak intermolecular  $C-H\cdots S$  hydrogen bonds (Table 1) link the molecules into spiral chains running along the *c* axis (Fig. 2). The relatively short distance  $Cl3\cdots Cl5^{iii}$  [symmetry code: (iii) 1 + x, *y*, *z*] of 3.364 (3) Å [ $Cl5-Cl3\cdots Cl5^{iii} = 158.1$  (3)°] suggests the existence of intermolecular  $Cl\cdots Cl$  interactions (Matsumoto *et al.*, 2002).



#### Figure 1

The molecular structure of (I), showing the atom-labelling scheme and displacement ellipsoids at the 50% probability level. Dashed lines indicate the intramolecular hydrogen bonds.

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## Experimental

The title compound was prepared according to the literature method of Zhang & Lin (1992). Single crystals suitable for X-ray analysis were obtained by slow evaporation of a solution in N,N-dimethyl-formamide at 293 K.

#### Crystal data

 $\begin{array}{l} C_{15}H_9Cl_5N_2O_2S\\ M_r=458.55\\ Tetragonal, P4_3\\ a=7.9580\ (11)\ \text{\AA}\\ c=29.249\ (6)\ \text{\AA}\\ V=1852.3\ (5)\ \text{\AA}^3\\ Z=4 \end{array}$ 

#### Data collection

Enraf-Nonius CAD-4 diffractometer  $\omega/2\theta$  scans Absorption correction:  $\psi$  scan (North *et al.*, 1968)  $T_{\min} = 0.713, T_{\max} = 0.839$ 3624 measured reflections

#### Refinement

Refinement on  $F^2$   $R[F^2 > 2\sigma(F^2)] = 0.057$   $wR(F^2) = 0.145$  S = 0.993624 reflections 226 parameters H-atom parameters constrained  $D_x = 1.644 \text{ Mg m}^{-3}$ Mo K\alpha radiation  $\mu = 0.91 \text{ mm}^{-1}$ T = 293 (2) K Prism, colourless 0.40 \times 0.20 \times 0.20 mm

3624 independent reflections 2336 reflections with  $I > 2\sigma(I)$  $\theta_{\text{max}} = 26.0^{\circ}$ 3 standard reflections every 200 reflections intensity decay: none

 $w = 1/[\sigma^{2}(F_{o}^{2}) + (0.07P)^{2}]$ where  $P = (F_{o}^{2} + 2F_{c}^{2})/3$  $(\Delta/\sigma)_{max} = 0.24 \text{ e} \text{ Å}^{-3}$  $\Delta\rho_{min} = -0.34 \text{ e} \text{ Å}^{-3}$ Absolute structure: Flack (1983), 1771 Friedel pairs Flack parameter: -0.01 (12)



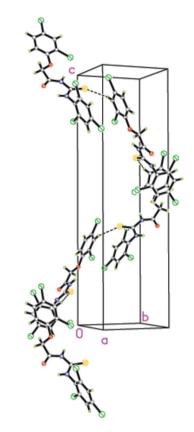
Hydrogen-bond geometry (Å, °).

$D - H \cdot \cdot \cdot A$	$D-\mathrm{H}$	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdots A$
N1-H1A····O1	0.86	2.05	2.516 (7)	113
$N2-H2A\cdots O2$	0.86	1.99	2.675 (7)	136
$C1 - H1B \cdot \cdot \cdot O2^{i}$	0.93	2.57	3.326 (9)	139
$C5-H5A\cdots S1^{ii}$	0.93	2.87	3.716 (7)	152

Symmetry codes: (i)  $y, -x + 2, z + \frac{1}{4}$ ; (ii)  $y + 1, -x + 1, z + \frac{1}{4}$ .

All H atoms were positioned geometrically (C-H = 0.93–0.97 Å; N-H = 0.86 Å) and refined as riding with  $U_{iso}(H) = 1.2U_{eq}(C,N)$ .

Data collection: *CAD-4 Software* (Enraf–Nonius, 1989); cell refinement: *CAD-4 Software*; data reduction: *XCAD4* (Harms & Wocadlo, 1995); program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *SHELXTL* (Bruker, 1998); software used to prepare material for publication: *SHELXTL*.



#### Figure 2

A portion of the crystal packing showing the spiral hydrogen-bonded (dashed lines) chain in (I).

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