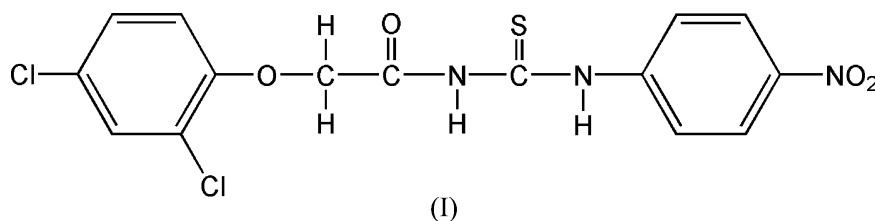


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Key indicators

Single-crystal X-ray study
 $T = 293$ K
Mean $\sigma(\text{C}-\text{C}) = 0.006$ Å
 R factor = 0.064
 wR factor = 0.170
Data-to-parameter ratio = 14.7For details of how these key indicators were
automatically derived from the article, see
<http://journals.iucr.org/e>.*N*-(2,4-Dichlorophenoxyacetyl)-*N'*-(4-nitrophenyl)thioureaIn the crystal structure of the title compound, $\text{C}_{15}\text{H}_{11}\text{Cl}_2\text{N}_3\text{O}_4\text{S}$, there are two intramolecular $\text{N}-\text{H}\cdots\text{O}$ and one $\text{C}-\text{H}\cdots\text{S}$ hydrogen bonds and a weak intermolecular $\text{C}-\text{H}\cdots\text{O}$ hydrogen bond.Received 28 June 2006
Accepted 17 August 2006

Comment

The title compound, (I), belongs to the class of aroylthiourea compounds, which have extensive biological properties, such as antiviral, herbicidal, pesticidal and plant-growth-regulating activity (Xu *et al.*, 2003; Sun *et al.*, 2006; Du *et al.*, 2002). Its structure is described here as part of our work studying the relationship between structure and biological activity.A view of the molecular structure of (I) is given in Fig. 1, and selected geometric parameters are listed in Table 1. The $\text{C9}-\text{S}$, $\text{C9}-\text{N1}$ and $\text{C9}-\text{N2}$ bond lengths are 1.644 (5), 1.395 (5) and 1.350 (5) Å, respectively, comparable with those in *N*-(4-methylbenzoyl)-*N'*-(4-nitrophenyl)thiourea [1.649 (2), 1.372 (3) and 1.339 (3) Å, respectively; Yusof *et al.*, 2006].The crystal structure of (I) contains two intramolecular $\text{N}-\text{H}\cdots\text{O}$ and one $\text{C}-\text{H}\cdots\text{S}$ hydrogen bonds, and a weak intermolecular $\text{C}-\text{H}\cdots\text{O}$ hydrogen bond. The intramolecular interactions form a five-membered $\text{N1}/\text{C8}/\text{C7}/\text{O1}/\text{H1A}$ ring and two six-membered rings, $\text{C8}/\text{N1}/\text{C9}/\text{C15}/\text{H15A}$ and $\text{O2}/\text{C8}/\text{N1}/\text{C9}/\text{N2}/\text{H2A}$.

Experimental

The title compound was prepared according to the literature method of Zhang *et al.* (1992). Single crystals suitable for X-ray analysis were obtained by slow evaporation of a dimethylformamide solution at 293 K.

Crystal data

 $\text{C}_{15}\text{H}_{11}\text{Cl}_2\text{N}_3\text{O}_4\text{S}$
 $M_r = 400.23$
Monoclinic, $C2/c$
 $a = 23.315$ (5) Å
 $b = 8.3190$ (17) Å
 $c = 18.452$ (4) Å
 $\beta = 109.18$ (3)°
 $V = 3380.2$ (14) Å³ $Z = 8$
 $D_x = 1.573$ Mg m⁻³
Mo $K\alpha$ radiation
 $\mu = 0.53$ mm⁻¹
 $T = 293$ (2) K
Prism, yellow
 $0.40 \times 0.20 \times 0.20$ mm

Data collection

Enraf–Nonius CAD-4
diffractometer
 $\omega/2\theta$ scans
Absorption correction: ψ scan
(*XCAD4*; Harms & Wocadlo,
1995)
 $T_{\min} = 0.815$, $T_{\max} = 0.901$
3400 measured reflections

3316 independent reflections
1943 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.040$
 $\theta_{\max} = 26.0^\circ$
3 standard reflections
every 200 reflections
intensity decay: none

Refinement

Refinement on F^2
 $R[F^2 > 2\sigma(F^2)] = 0.064$
 $wR(F^2) = 0.170$
 $S = 1.08$
3316 reflections
226 parameters
H-atom parameters constrained

$w = 1/[\sigma^2(F_o^2) + (0.065P)^2 + 3.2P]$
where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} < 0.001$
 $\Delta\rho_{\max} = 0.35 \text{ e } \text{\AA}^{-3}$
 $\Delta\rho_{\min} = -0.32 \text{ e } \text{\AA}^{-3}$

Table 1

Selected geometric parameters (\AA , $^\circ$).

S—C9	1.644 (5)	N2—C9	1.350 (5)
O1—C4	1.349 (5)	N2—C10	1.412 (5)
O1—C7	1.413 (5)	C7—C8	1.509 (6)
N1—C8	1.355 (5)	C10—C15	1.371 (6)
N1—C9	1.395 (5)	C10—C11	1.381 (5)
C4—O1—C7	119.5 (3)	N1—C8—C7	115.8 (4)
C8—N1—C9	130.2 (4)	N2—C9—N1	113.0 (4)
C9—N2—C10	131.2 (4)	N2—C9—S	129.5 (3)
O1—C7—C8	108.8 (4)	N1—C9—S	117.5 (3)
O2—C8—N1	125.4 (4)	C15—C10—N2	125.6 (4)
O2—C8—C7	118.7 (4)		

Table 2

Hydrogen-bond geometry (\AA , $^\circ$).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
N1—H1A \cdots O1	0.86	2.09	2.559 (5)	114
N2—H2A \cdots O2	0.86	1.95	2.678 (5)	142
C11—H11A \cdots O2 ¹	0.93	2.45	3.305 (5)	152
C15—H15A \cdots S	0.93	2.56	3.225 (5)	129

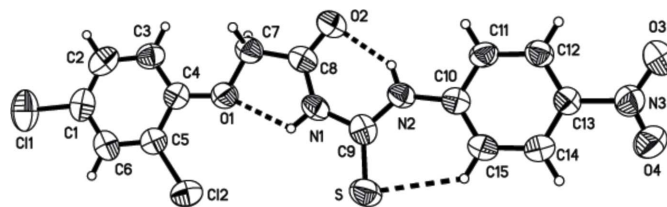
Symmetry code: (i) $-x + 1, y, -z - \frac{1}{2}$.

Figure 1

View of the molecular structure of (I), showing the atom-labelling scheme. Displacement ellipsoids are drawn at the 50% probability level. Dashed lines indicate hydrogen bonds.

All H atoms were positioned geometrically ($N-H = 0.86 \text{ \AA}$ and $C-H = 0.93$ or 0.97 \AA) and refined using a riding model, with $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C}, \text{N})$.

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*.

We are grateful to H.-Q. Wang for the crystal data collection.

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