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Key indicators

Single-crystal X-ray study T = 292 KMean $\sigma(\text{C}-\text{C}) = 0.003 \text{ Å}$ R factor = 0.042 wR factor = 0.122 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.

2-(2,4-Dichlorophenoxy)-*N*-[5-(3-pyridyl)-1,3,4-thiadiazol-2-yl]acetamide

In the title compound, $C_{15}H_{10}Cl_2N_4O_2S$, the three rings are nearly coplanar. In the crystal structure, intermolecular N– $H \cdot \cdot \cdot N$, C– $H \cdot \cdot \cdot Cl$ and C– $H \cdot \cdot \cdot N$ hydrogen-bond interactions are observed. Intermolecular π - π stacking interactions are also present. Received 15 June 2005 Accepted 17 June 2005 Online 24 June 2005

Comment

Thiadiazole derivatives exhibit many important bioactivities (Wang *et al.*, 2004; Castro *et al.*, 1996). The title compound, (I), is an example of this class.



The molecule of (I) (Fig. 1) is essentially planar; the dihedral angles formed by the thiadiazole ring with the benzene and pyridine planes are 0.3 (1) and 10.3 (1)°, respectively. All bond lengths and angles in (I) are as expected (Table 1).

In the crystal structure, intermolecular N-H···N, C-H···Cl and C-H···N interactions are observed, as shown in Fig. 2. Also present are π - π stacking interactions (Fig. 3) between the thiadiazole and benzene rings. The interplanar spacing is 3.497 (2) Å, the centroid-to-centroid separation is 3.631 (2) Å and the centroid offset is 0.997 (2) Å.

Experimental

2-Amino-5-(3-pyridyl)-1,3,4-thiadiazole (0.88 g, 4.0 mmol) was prepared according to the reported procedure of Song *et al.* (2005). It was then treated with 2,4-dichlorophenoxyethyl acid chloride (1.08 g, 4.5 mmol), which was readily available in 90% yield by refluxing 2,4-dichlorophenoxyethyl acid (obtained commercially and used without



Figure 1

© 2005 International Union of Crystallography Printed in Great Britain – all rights reserved A view of the molecule of (I), showing the atom-labelling scheme. Displacement ellipsoids are drawn at the 50% probability level.



Figure 2

A partial packing diagram for (I) [symmetry codes: (a) 1 - x, -y, -z; (b) $1 - x, \frac{1}{2} + y, \frac{1}{2} - z$]. Hydrogen-bonding interactions are indicated by dashed lines.

further purification) with an excess of thionyl chloride. The title compound, (I), was isolated in 73% yield. Crystals suitable for X-ray analysis were obtained by slow evaporation of a methanol-dimethylformamide (1:3) solution at room temperature (m.p. 528–529 K). Elemental analysis: analysis calculated for $C_{15}H_{10}Cl_2N_4O_2S$: C 47.26, H 2.64, N 14.70%; found: C 47.37, H 2.82, N 14.55%.

Crystal data

$C_{15}H_{10}Cl_2N_4O_2S$	$D_x = 1.601 \text{ Mg m}^{-3}$
$M_r = 381.23$	Mo $K\alpha$ radiation
Monoclinic, $P2_1/c$	Cell parameters from 2934
a = 7.6550 (7) Å	reflections
b = 9.0221 (8) Å	$\theta = 2.4-27.7^{\circ}$
c = 22.927 (2) Å	$\mu = 0.56 \text{ mm}^{-1}$
$\beta = 92.811 \ (2)^{\circ}$	T = 292 (2) K
V = 1581.6 (2) Å ³	Block, colourless
Z = 4	$0.30 \times 0.20 \times 0.20 \ \text{mm}$
Data collection	

Bruker SMART CCD area-detector diffractometer φ and ω scans Absorption correction: multi-scan

(SADABS; Sheldrick, 1996) $T_{min} = 0.850, T_{max} = 0.896$ 8360 measured reflections 3100 independent reflections 2641 reflections with $I > 2\sigma(I)$ $R_{int} = 0.024$ $\theta_{max} = 26.0^{\circ}$ $h = -9 \rightarrow 6$ $k = -10 \rightarrow 11$ $l = -28 \rightarrow 28$



Figure 3

Part of the crystal structure of (I), showing the π - π stacking interactions. [Symmetry codes: (a) x, y – 1, z; (b) x, y + 1, z.]

Refinement on F^2	$w = 1/[\sigma^2(F_0^2) + (0.0684P)]$
$R[F^2 > 2\sigma(F^2)] = 0.042$	+ 0.38P]
$wR(F^2) = 0.122$	where $P = (F_0^2 + 2F_c^2)/3$
S = 1.08	$(\Delta/\sigma)_{\rm max} = 0.001$
3100 reflections	$\Delta \rho_{\rm max} = 0.39 \ {\rm e} \ {\rm \AA}^{-3}$
217 parameters	$\Delta \rho_{\rm min} = -0.21 \text{ e } \text{\AA}^{-3}$
H-atom parameters constrained	

Table 1

Selected geometric parameters (Å, °).

C6-O1	1.368 (3)	C9-N1	1.368 (3)
C7-C8	1.509 (3)	C10-N3	1.302 (3)
C8-O2	1.207 (3)	C10-C11	1.465 (3)
C8-N1	1.368 (3)	C15-N4	1.328 (3)
C9-N2	1.298 (3)	C14-N4	1.338 (3)
O2-C8-N1	123.3 (2)	N3-C10-C11	123.07 (19)
O2-C8-C7	127.1 (2)	C8-N1-C9	125.25 (19)
N1-C8-C7	109.58 (18)	C6-O1-C7	114.39 (17)
01-C7-C8-O2	9.0 (4)	O2-C8-N1-C9	2.0 (4)
O1-C7-C8-N1	-172.60(19)	S1-C9-N1-C8	-5.9(3)
S1-C10-C11-C15	-169.10 (18)	C5-C6-O1-C7	-2.3 (3)

Table 2

Hydrogen-bond geometry (Å, °).

$D - H \cdot \cdot \cdot A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdot \cdot \cdot A$
C14-H14···Cl1 ⁱ	0.93	2.81	3.552 (3)	138
$C7 - H7A \cdot \cdot \cdot N2^{ii}$	0.97	2.42	3.230 (3)	141
$N1 - H1 \cdots N3^{ii}$	0.86	2.09	2.951 (3)	176
		. 11		

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

All H atoms were placed in geometrically idealized positions and constrained to ride on their parent atoms, with C-H distances of 0.93 Å for phenyl and pyridyl H and 0.97 Å for methylene H, and N-H distances of 0.86 Å, and with $U_{iso}(H) = 1.2U_{eq}(C,N)$.

Data collection: *SMART* (Bruker, 2001); cell refinement: *SAINT* (Bruker, 2001); data reduction: *SAINT*; structure solution: *SHELXS97* (Sheldrick, 1997); structure refinement: *SHELXL97* (Sheldrick, 1997); molecular graphics: *SHELXTL* (Sheldrick, 2001); software used to prepare material for publication: *SHELXTL*.

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