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Masoumeh Tabatabaee,^a* Mitra Ghassemzadeh,^b Ali Reza Dehghan,^a Hamid Reza Khavasi^c and Majid M. Heravi^d

^aDepartment of Chemistry, Islamic Azad University, Yazd Branch, Yazd, Iran, ^bChemistry & Chemical Engineering, Research Center of Iran, Tehran, Iran, ^cDepartment of Chemistry, Shahid Beheshti University, Evin, Tehran, Iran, and ^dSchool of Sciences, Azahra University, Tehran, Iran

Correspondence e-mail: tabatabaee45m@yahoo.com

Key indicators

Single-crystal X-ray study T = 150 KMean σ (C–C) = 0.002 Å R factor = 0.029 wR factor = 0.067 Data-to-parameter ratio = 17.1

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

Ethoxycarbonylmethyl 3-(2-chlorobenzylidene)dithiocarbazate

The title compound, $C_{12}H_{13}ClN_2O_2S_2$, is a new Schiff base derived from ring opening of *N*-aminorhodanine. It is prepared by reaction of *N*-aminorhodanine and 2-chlorobenzaldehyde in an ethanol solution in the presence of HCl. In the solid state, intermolecular $N-H \cdots O$ hydrogen bonds link the molecules into chains.

Comment

During the last few decades, there has been considerable interest in the chemistry of Schiff base compounds (Dubey & Vaid, 1991). Schiff bases have been extensively studied for various reasons, one of which is their biological activity (Todeschini et al., 1998; Demirbas, 2004; Rando et al., 2002). Schiff bases containing different donor atoms also find use in analytical applications (Galic et al., 2001; Wyrzykiewicz & Prukah, 1998; Reddy & Lirgappa, 1994). Recently, we have reported the syntheses of some Schiff bases derived from 4amino-6-methyl-1,2,4-triazine-3-thione-5(2H)-one (AMTTO; Tabatabaee et al., 2006) and 4-amino-5-methyl-1,2,4-triazole-3(4H)-thione (AMTT), and their use as ligands in Ag^I and Cu^I complexes (Ghassemzadeh et al. 2004, 2005, 2006). We report here a new Schiff base, (I), derived from ring opening of Naminorhodanine, prepared by its reaction with 2-chlorobenzaldehyde in ethanol solution in the presence of HCl.



The molecular structure of (I) is shown in Fig. 1. Bond lengths and angles are unexceptional. In the solid state, intermolecular $N-H\cdots O$ hydrogen bonds (Table 1) link the molecules into chains.



Figure 1

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The molecular structure of (I), showing displacement ellipsoids drawn at the 50% probability level. H atoms are shown as spheres of arbitrary radius.

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Experimental

A solution of *N*-aminorhodanine (4 mmol) in EtOH (20 ml) was treated with 2-chlorobenzaldehyde (6 mmol) and the resulting mixture was acidified with 37% hydrochloric acid (0.2 ml). The reaction mixture was refluxed for 5 h. The solid residue was filtered, washed with cold ethanol (10 ml) and recrystallized from EtOH (yield 89%).

Z = 4

 $D_x = 1.468 \text{ Mg m}^{-3}$

 $0.20 \times 0.15 \times 0.10 \ \text{mm}$

10851 measured reflections

3029 independent reflections

2808 reflections with $I > 2\sigma(I)$

Mo $K\alpha$ radiation

 $\mu = 0.56 \text{ mm}^{-1}$ T = 150 (2) K Prism, vellow

 $R_{\rm int} = 0.028$

 $\theta_{\rm max} = 26.8^\circ$

Crystal data

$C_{12}H_{13}CIN_2O_2S_2$
$M_r = 316.81$
Monoclinic, $P2_1/c$
a = 10.5613 (18) Å
b = 16.856 (3) Å
c = 8.1013 (14) Å
$\beta = 96.409 \ (14)^{\circ}$
V = 1433.2 (4) Å ³

Data collection

Stoe IPDS-II diffractometer φ scans Absorption correction: numerical (X-SHAPE; Stoe & Cie, 2005) $T_{\min} = 0.905, T_{\max} = 0.946$

Refinement

Refinement on F^2	$w = 1/[\sigma^2(F_0^2) + (0.0242P)^2]$	Ghassemzadeh, M.
$R[F^2 > 2\sigma(F^2)] = 0.029$	+ 0.6651P]	A. & Neumüller
$wR(F^2) = 0.067$	where $P = (F_0^2 + 2F_c^2)/3$	Ghassemzadeh, M.
S = 1.14	$(\Delta/\sigma)_{\rm max} = 0.001$	Anorg. Allg. Che
3029 reflections	$\Delta \rho_{\rm max} = 0.27 \text{ e} \text{ \AA}^{-3}$	Rando, D. G., Sato,
177 parameters	$\Delta \rho_{\rm min} = -0.17 \text{ e } \text{\AA}^{-3}$	T., Ferreira, F. I.
H atoms treated by a mixture of		Reddy, H. & Lirga
independent and constrained		Sheldrick, G. M.
refinement		Göttingen, Gern
		Stoe & Cie (2000

Table 1

Hydrogen-bond geometry (Å, °).

$D - H \cdots A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdots A$
$N1 - H1 \cdots O2^i$	0.85 (2)	2.17 (2)	2.9870 (18)	162.0 (19)

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

H atoms were positioned geometrically, with C–H = 0.95 Å for aromatic, C–H = 0.99 Å for methylene and C–H = 0.98 Å for methyl H atoms, and were allowed to ride during subsequent refinement, with $U_{iso}(H) = 1.2U_{eq}(C)$ or $1.5U_{eq}(methyl C)$. The methyl group was allowed to rotate about its local threefold axis. The H atom of the NH group was located in a difference Fourier map and refined freely with an isotropic displacement parameter.

Data collection: X-AREA (Stoe & Cie, 2005); cell refinement: X-RED32 (Stoe & Cie, 2005); data reduction: X-RED32; program(s) used to solve structure: SHELXS97 (Sheldrick, 1997); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: X-STEP32 (Stoe & Cie, 2000); software used to prepare material for publication: SHELXL97.

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