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Key indicators

Single-crystal X-ray study T = 294 KMean $\sigma(\text{C}-\text{C}) = 0.003 \text{ Å}$ R factor = 0.041 wR factor = 0.113 Data-to-parameter ratio = 15.3

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

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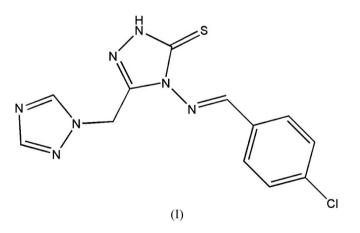
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4-(4-Chlorobenzylideneamino)-3-(1*H*-1,2,4-triazol-1-ylmethyl)-1*H*-1,2,4-triazole-5(4*H*)-thione

In the title compound, $C_{12}H_{10}CIN_7S$, the dihedral angles made by the plane of the thione-substituted triazole ring with the planes of the other triazole ring and the benzene ring are 73.57 (3) and 46.65 (2)°, respectively. Inter- and intramolcular hydrogen bonds and π - π stacking interactions stabilize the structure. Received 15 September 2005 Accepted 20 September 2005 Online 24 September 2005

Comment

Recently, compounds containing a 1*H*-1,2,4-triazole group have attracted much interest because compounds containing this ring system are well known as efficient fungicides in pesticides and they exhibit good plant-growth regulatory activity for a wide variety of crops (Xu *et al.*, 2002). In addition, amine- and thione-substituted triazoles have been studied as anti-inflammatory and antimicrobial agents (Eweiss *et al.*,1986; Awad *et al.*, 1991). In a search for new triazole compounds with better biological activity, the title compound, (I), was synthesized. We report here the crystal structure of (I).



Bond lengths and angles in the thione-substituted triazole ring and the other triazole ring (Table 1) are in agreement with the values quotes in previous reports (Li *et al.*, 2005; Xu *et al.*, 2005). The molecule exists in the thione tautomeric form, with an S=C distance of 1.662 (2) Å, which indicates substantial double-bond character for this bond (Allen *et al.*, 1987). The planes C10–C12/N5/N6/N7 and C1–C7/N1/C11 make angles of 73.57 (3) and 46.65 (2)°, respectively, with the thione-substituted triazole plane C8/C9/N2/N3/N4/S1. The crystal structure of (I) is stabilized by weak intra- and intermolecular hydrogen bonds and π - π stacking interactions. The π - π stacking interactions involve the thione-substituted triazole ring (C8/C9/N2/ N3/N4/S1) *R*1 and the benzene ring (C2–C7) *R*2. The distance between the centroids of rings *R*1 and *R*2 (at 1 - x, $\frac{1}{2} + y$, $\frac{1}{2} - z$) is 3.532 (3) Å.

Experimental

A mixture of 4-amino-3-(1,2,4-triazol-1-yl)-1*H*-1,2,4-triazole-5(4*H*)thione (0.02 mol) and 4-chlorobenzaldehyde (0.02 mol) was refluxed at 391 K for 15–20 min in glacial acetic acid. The mixture was then filtered and crystallized from ethanol to afford the title compound (5.87 g, yield 92%). Single crystals suitable for X-ray measurements were obtained by recrystallization from ethanol at room temperature.

 $D_x = 1.464 \text{ Mg m}^{-3}$

Mo $K\alpha$ radiation Cell parameters from 2083

reflections $\theta = 2.3-25.2^{\circ}$ $\mu = 0.41 \text{ mm}^{-1}$

T = 294 (2) K

Block, yellow $0.22 \times 0.20 \times 0.18 \text{ mm}$

 $R_{\rm int} = 0.029$

 $\theta_{\rm max} = 26.4^{\circ}$

 $h = -8 \rightarrow 13$

 $k = -9 \rightarrow 8$

 $l = -22 \rightarrow 22$

2972 independent reflections

1855 reflections with $I > 2\sigma(I)$

Crystal data

C ₁₂ H ₁₀ ClN ₇ S
$M_r = 319.78$
Monoclinic, $P2_1/c$
a = 11.090 (2) Å
b = 7.4421 (15)Å
c = 17.734 (4) Å
$\beta = 97.693 \ (3)^{\circ}$
$V = 1450.6 (5) \text{ Å}^3$
Z = 4

Data collection

Bruker SMART CCD area detector diffractometer φ and ω scans Absorption correction: multi-scan (*SADABS*; Sheldrick, 1996) $T_{\rm min} = 0.910, T_{\rm max} = 0.929$ 7928 measured reflections

Refinement

-	
Refinement on F^2	$w = 1/[\sigma^2(F_0^2) + (0.05P)^2]$
$R[F^2 > 2\sigma(F^2)] = 0.041$	+ 0.2972P]
$wR(F^2) = 0.113$	where $P = (F_0^2 + 2F_c^2)/3$
S = 1.01	$(\Delta/\sigma)_{\rm max} = 0.001$
2972 reflections	$\Delta \rho_{\rm max} = 0.16 \ {\rm e} \ {\rm \AA}^{-3}$
194 parameters	$\Delta \rho_{\rm min} = -0.27 \text{ e } \text{\AA}^{-3}$
H atoms treated by a mixture of	
independent and constrained	
refinement	

Table 1

Selected geometric parameters (Å, $^{\circ}$).

S1-C8	1.662 (2)	N3-N4	1.376 (2)
N1-N2	1.395 (2)	N5-N6	1.354 (3)
C1-N1-N2	116.02 (18)	N5-C10-C9	112.72 (18)

Table 2

Hydrogen-bond geometry (Å, °).

$D - H \cdot \cdot \cdot A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - H \cdots A$
N4-H4A···N7 ⁱ	0.83 (2)	2.01 (3)	2.830 (3)	169 (2)
$C4-H4\cdots S1^{ii}$	0.93	2.82	3.517 (2)	133
$C1 - H1 \cdots S1$	0.93	2.74	3.246 (2)	115

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

All H atoms were placed in calculated positions, with C-H = 0.93–0.97 Å, and refined using a riding model, with $U_{\rm iso}({\rm H}) = 1.2U_{\rm eq}({\rm C})$.

Data collection: *SMART* (Bruker, 1998); cell refinement: *SAINT* (Bruker, 1999); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *SHELXTL* (Bruker, 1999); software used to prepare material for publication: *SHELXTL*.

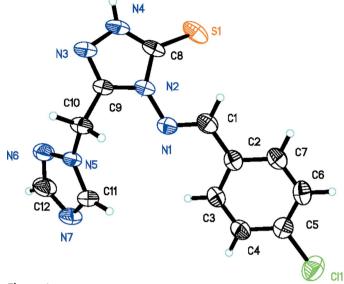


Figure 1

View of the title compound (I), with displacement ellipsoids drawn at the 40% probability level.

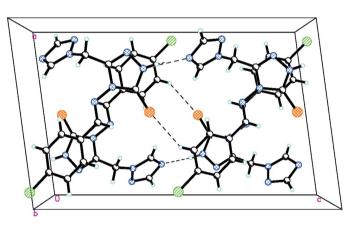


Figure 2

The packing of the title compound, viewed down the b axis. Hydrogen bonds are shown as dashed lines.

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