Acta Crystallographica Section E Structure Reports Online

ISSN 1600-5368

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Key indicators

Single-crystal X-ray study T = 100 KMean σ (C–C) = 0.002 Å R factor = 0.038 wR factor = 0.083 Data-to-parameter ratio = 20.4

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3-Benzyl-4-(4-methylphenyl)-1*H*-1,2,4-triazole-5(4*H*)-thione

The title compound, $C_{16}H_{15}N_3S$, displays the usual geometrical parameters of 1,2,4-triazole derivatives. The occurrence of $N-H\cdots S$ hydrogen-bonding interactions results in the formation of dimers.

Received 12 October 2004 Accepted 3 November 2004 Online 13 November 2004

Comment

Derivatives of 1,2,4-triazole are known to exhibit antiinflammatory (Unangst et al., 1992; Mullican et al., 1993), antiviral, analgesic (Sughen & Yoloye, 1978), antimicrobial (Shams El-Dine & Hazzaa, 1974; Misato et al., 1977; Cansız et al., 2001), anticonvulsant (Stillings et al., 1986) and antidepressant activity (Kane et al., 1988), the latter being usually explored by the forced swim test (Porsolt et al., 1977; Vamvakides, 1990). Among the pharmacological profiles of 1,2,4-triazoles, their antimicrobial, anticonvulsant and antidepressant properties seem to be the best documented. Derivatives of 4,5-disubstituted 1,2,4-triazole are known to be synthesized by intramolecular cyclization of 1,4 disubstituted thiosemicarbazides (Zamani et al., 2003; Cansız et al., 2004; Koparır et al., 2004). In addition, there are some studies on the electronic structures and thiol-thione tautomeric equilibrium of heterocyclic thione derivatives (Aydoğan et al., 2002; Charistos et al., 1994). In the present study, the synthesis and structural characterization of 3-benzyl-4-(4-methylphenyl)-1H-1,2,4-triazole-5(4H)-thione, (III), are reported.



The 1,2,4-triazole ring is planar, the maximum deviation from planarity being 0.002 (1) Å for atom N2 (Fig. 1). The tolyl ring is twisted by 67.38 (5)° around the N1–C10 bond with respect to the triazole ring, whereas the phenyl and triazole rings make a dihedral angle of 68.38 (5)°.

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ORTEP-3 view (Farrugia, 1997) of the title compound, showing 50% probability displacement ellipsoids and the atom-numbering scheme.



Figure 2

A packing diagram of (III), illustrating the formation of dimers through N-H···S hydrogen bonding (dashed lines).

All bond lengths and angles have values comparable with those reported in the literature for related structures (Öztürk et al., 2004; Akkurt et al., 2004).

In the crystal structure, the occurrence of intermolecular N-H···S hydrogen bonding (Table 2) results in the formation of dimers (Fig. 2).

Experimental

The title compound, (III), was synthesized by the reaction of 1-isothiocyanato-4-methylbenzene and 2-phenylacetohydrazide, (I), through the intermediate N-(4-methylphenyl)-2-(phenylacetyl)hydrazinecarbothioamide, (II). Base-catalysed intramolecular dehydrative cyclization of this intermediate furnished the thione in good yield. For the synthesis of (II), a mixture of (I) (0.01 mol) and the appropriate 1-isothiocyanato-4-methylbenzene (0.01 mol) in absolute ethanole (100 ml) were refluxed for 7 h. The solid material obtained on cooling was filtered off, washed with diethyl ether, dried and crystallized from methanol (yield 88%; m.p. 433–434 K). IR ν (cm⁻¹): 3450-3329 (N-H), 1678 (C=O), 1250 (C=S). ¹H NMR: δ 2.28 (s, 3H, CH₃), 3.53 (s, 2H, CH₂), 7.30-7.45 (m, 9H, Ar. H), 8.17-8.24 (br, 1H, -NH-Ar); 10.13–10.15 (br, 2H, 2 × NH); analysis calculated for C₁₆H₁₇N₃OS (299): C 64.19, H 5.72, N 14.03, S 10.71%; found C 64.11, H 5.69, N 14.01, S 10.59%.

For the synthesis of (III), a stirred mixture of (II) (1 mmol) and sodium hydroxide (40 mg, 1 mmol, as a 2 N solution) was refluxed for 4 h. After cooling, the solution was acidified with hydrochloric acid and the precipitate was filtered off. The precipitate was then crystallized from a mixture of methanol-dioxane (yield: 71%; m.p.: 455-456 K). IR ν (cm⁻¹): 3358–3290 (NH), 2550 (SH), 1605 (C=N), 1538, 1262, 1050, 950 (N–C=S, amide I, II, III and IV bands); ¹H NMR δ: 2.42 (s, 3H, CH₃), 3.81 (s, 2H, CH₂), 6.97–7.29 (m, 9H, Ar. H), 12.91 (s, 1H, SH/NH); analysis calculated for C₁₆H₁₅N₃S (312): C 68.30, H 5.37, N 14.93, S 11.40%; found C 63.38, H 5.37, N 15.00, S 11.47%.

Crystal data

	2
$C_{16}H_{15}N_3S$	$D_x = 1.320 \text{ Mg m}^{-3}$
$M_r = 281.38$	Mo $K\alpha$ radiation
Monoclinic, $P2_1/n$	Cell parameters from 119
a = 12.3833 (12) Å	reflections
b = 7.2297 (5) Å	$\theta = 6-20^{\circ}$
c = 15.8926 (15) Å	$\mu = 0.22 \text{ mm}^{-1}$
$\beta = 95.488 \ (7)^{\circ}$	$T = 100 { m K}$
$V = 1416.3 (2) \text{ Å}^3$	Prism, colorless
Z = 4	$0.27 \times 0.23 \times 0.18 \text{ mm}$

Data collection

Bruker–Nonius KappaCCD	3712 independent reflections
diffractometer	2826 reflections with $I > 2\sigma(I)$
ω – φ scans	$R_{\rm int} = 0.047$
Absorption correction: multi-scan	$\theta_{\rm max} = 29.0^{\circ}$
(SADABS; Sheldrick, 2002)	$h = -16 \rightarrow 16$
$T_{\min} = 0.929, \ T_{\max} = 0.962$	$k = -9 \rightarrow 9$
21959 measured reflections	$l = -21 \rightarrow 21$

Refinement

Refinement on F^2 H-atom parameters constrained $R[F^2 > 2\sigma(F^2)] = 0.038$ $w = 1/[\sigma^2(F_o^2) + (0.0358P)^2]$ $wR(F^2) = 0.084$ + 0.5394P] where $P = (F_o^2 + 2F_c^2)/3$ S = 1.023712 reflections $(\Delta/\sigma)_{\rm max} = 0.001$ $\Delta \rho_{\rm max} = 0.29 \ {\rm e} \ {\rm \AA}^{-3}$ 182 parameters $\Delta \rho_{\rm min} = -0.26 \ {\rm e} \ {\rm \AA}^{-3}$

Table 1

Selected geometric parameters (Å, °).

1.6853 (13)	N1-C10	1.4421 (16)
1.3750 (17)	N2-C1	1.3417 (16)
1.3850 (16)	N2-N3	1.3805 (15)
107.80 (10)	C1-N2-N3	113.44 (11)
124.41 (11)	C2-N3-N2	104.03 (10)
127.72 (11)		. ,
	1.6853 (13) 1.3750 (17) 1.3850 (16) 107.80 (10) 124.41 (11) 127.72 (11)	1.6853 (13) N1-C10 1.3750 (17) N2-C1 1.3850 (16) N2-N3 107.80 (10) C1-N2-N3 124.41 (11) C2-N3-N2 127.72 (11) C1

organic papers

Table 2Hydrogen-bonding geometry (Å, $^{\circ}$).

$D - H \cdots A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdots A$
$N2-H2\cdots S1^i$	0.86	2.42	3.2772 (12)	174

Symmetry code: (i) 1 - x, -y, 1 - z.

All the H atoms were located in difference maps, but they were idealized and treated as riding on their parent C or N atoms $[N-H = 0.86, C-H = 0.93-0.97 \text{ Å}; U_{iso}(H) = 1.2 \text{ or } 1.5 \text{ times } U_{eq}(\text{parent atom})].$

Data collection: *COLLECT* (Nonius, 1999); cell refinement: *EVALCCD* (Duisenberg *et al.*, 2003); data reduction: *EVALCCD*; program(s) used to solve structure: *SHELXS*97 (Sheldrick, 1997); program(s) used to refine structure: *SHELXL*97 (Sheldrick, 1997); molecular graphics: *ORTEP-3 for Windows* (Farrugia, 1997) and *PLATON* (Spek, 2003); software used to prepare material for publication: *WinGX* (Farrugia, 1999).

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