Acta Crystallographica Section E Structure Reports Online

ISSN 1600-5368

Derya Ugur,^a Ulrich Flörke,^b* Nevzat Külcü^a and Hakan Arslan^a

^aDepartment of Chemistry, Faculty of Arts and Science, Mersin University, Mersin, Turkey, and ^bDepartment Chemie, Fakultät für Naturwissenschaften, Universität Paderborn, Warburgerstrasse 100, D-33098 Paderborn, Germany

Correspondence e-mail: uf@chemie.uni-paderborn.de

Key indicators

Single-crystal X-ray study T = 173 K Mean σ (C–C) = 0.002 Å R factor = 0.032 wR factor = 0.090 Data-to-parameter ratio = 16.1

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

complexation capacity.

Comment

The centre of the molecule lies about a crystallographic inversion centre and thus the two thiourea moieties adopt an anti position. The planes of the benzoyl ring and the thiourea N₂CS group are almost perpendicular with an angle of 89.0 (1)°. The corresponding torsion angles are N2-C6- $C7-C8 = 160.4 (1)^{\circ}, C6-N2-C5-S1 = 108.3 (1)^{\circ} \text{ and } O1 C6-C7-C8 = 154.8 (1)^{\circ}$. Prominent bond lengths are in the expected range, viz. C5-S1 = 1.6660(13) Å, C6-O1 =1.2185 (16) Å, C5-N2 = 1.4173 (16) Å and C5-N1 =1.3214 (17) Å and compare well with the related distances of 1,1-diethyl-3-(4-methylbenzoyl)thiourea (Morales et al., 1997) or N-benzoyl-N'-methyl-N'-phenylthiourea (Shanmuga Sundara Raj et al., 1999). In the crystal structure (Fig. 2), molecules form dimers through strong intermolecular N2- $H2 \cdot \cdot \cdot S1(1 - x, -y, 2 - z)$ hydrogen bonds, with $H \cdot \cdot \cdot S =$ 2.37 Å and $N-H \cdot \cdot \cdot S = 172^{\circ}$. An additional intermolecular interaction is C2-H2B···O1 $(x-\frac{1}{2}, \frac{1}{2}-y, z-\frac{1}{2})$, with $H \cdots O = 2.33 \text{ Å}$ and $C - H \cdots O = 152^{\circ}$. An intramolecular hydrogen bond is formed by C2-H2A···N2, with H···N =

3-[4-(3,3-Diethylthioureidocarbonoyl)-

The title compound, C₁₈H₂₆N₄O₂S₂, lies about a crystal-

lographic inversion centre. The crystal packing shows inter-

molecular $C-H\cdots O$ and $N-H\cdots S$ interactions, the latter

The title compound, (I), is another example of our newly

synthesized thiourea derivatives which show interesting

(I)

benzoyl]-1,1-diethylthiourea

giving rise to the formation of dimers.



Figure 1

The molecular structure of the title compound. Displacement ellipsoids are drawn at the 50% probability level. [Symmetry code: (A) 1-x, -y, 3-z.]

Received 13 August 2003 Accepted 14 August 2003 Online 23 August 2003

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Figure 2

Packing diagram, viewed along [100]. Intermolecular hydrogen bonding is indicated by dashed lines.

2.30 Å and C-H···N = 105°. All these values are normalized for N-H = 1.03 Å and C-H = 1.08 Å.

Experimental

A solution of 2.5 mmol terephthalyl dichloride in 125 ml acetone was added to 5 mmol KSCN in 25 ml of acetone. The mixture was stirred for 30 min at 313 K and then cooled to room temperature. Afterwards a solution of 5 mmol diethylamine in 25 ml acetone was added dropwise with stirring, which was continued for 2 h. The yellowish precipitate was recrystallized from dichloromethane–ethanol (1:1).

Crystal data

$C_{18}H_{26}N_4O_2S_2$	$D_x = 1.332 \text{ Mg m}^{-3}$	
$M_r = 394.55$	Mo $K\alpha$ radiation	
Monoclinic, $P2_1/n$	Cell parameters from 3822	
a = 6.7980(7) Å	reflections	
b = 15.2735 (16) Å	$\theta = 2.5 - 28.3^{\circ}$	
c = 10.0275 (10) Å	$\mu = 0.29 \text{ mm}^{-1}$	
$\beta = 109.168 \ (1)^{\circ}$	T = 173 (2) K	
$V = 983.43 (17) \text{ Å}^3$	Prism, colourless	
Z = 2	$0.32 \times 0.25 \times 0.12 \text{ mm}$	

Data collection

refinement

Bruker SMART CCD area-detector diffractometer	1991 independent reflections 1821 reflections with $I > 2\sigma(I)$
φ and ω scans	$R_{\rm int} = 0.028$
Absorption correction: multi-scan	$\theta_{\rm max} = 26.4^{\circ}$
(SADABS; Bruker, 2002)	$h = -8 \rightarrow 8$
$T_{\rm min} = 0.890, \ T_{\rm max} = 0.973$	$k = -19 \rightarrow 17$
5476 measured reflections	$l = -12 \rightarrow 12$
Refinement	
Refinement on F^2	$w = 1/[\sigma^2(F_o^2) + (0.0466P)^2]$
$R[F^2 > 2\sigma(F^2)] = 0.032$	+ 0.2868P]
$wR(F^2) = 0.090$	where $P = (F_o^2 + 2F_c^2)/3$
S = 1.06	$(\Delta/\sigma)_{\rm max} = 0.001$
1991 reflections	$\Delta \rho_{\rm max} = 0.29 \ {\rm e} \ {\rm \AA}^{-3}$
124 parameters	$\Delta \rho_{\rm min} = -0.17 \mathrm{e} \mathrm{\AA}^{-3}$
H atoms treated by a mixture of	
independent and constrained	

 Table 1

 Selected geometric parameters (Å, °).

\$1-C5	1.6660 (13)	N2-C6	1.3606 (17)
O1-C6	1.2185 (16)	N2-C5	1.4173 (16)
N1-C5	1.3214 (17)	C6-C7	1.4915 (17)
C6-N2-C5	119.59 (10)	O1-C6-N2	122.06 (12)
N1-C5-N2	115.88 (11)	O1-C6-C7	121.05 (12)
N1-C5-S1	125.10 (10)	N2-C6-C7	116.85 (11)
N2-C5-S1	119.02 (9)		

The H atom bonded to N2 was refined freely. Other H atoms were placed at calculated positions, riding on their attached C atoms (C– H = 0.95, 0.98 and 0.99 Å for aromatic, CH₂ and CH₃ H atoms, respectively), with isotropic displacement parameters $U_{\rm iso}(H) = 1.2U_{\rm eq}(C)$ or $1.5U_{\rm eq}(CH_3)$. CH₃ groups were allowed to rotate, but not to tip.

Data collection: *SMART* (Bruker, 2002); cell refinement: SMART; data reduction: *SAINT* (Bruker, 2002); program(s) used to solve structure: *SHELXTL* (Bruker, 2002); program(s) used to refine structure: *SHELXTL*; molecular graphics: *SHELXTL*; software used to prepare material for publication: *SHELXTL*.

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