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Key indicators

Single-crystal X-ray study T = 290 KMean $\sigma(C-C) = 0.003 \text{ Å}$ R factor = 0.054 wR factor = 0.127 Data-to-parameter ratio = 12.5

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

1,3-Bis(4-methoxyphenyl)thiourea

In the crystal structure of the title compound, $C_{15}H_{16}N_2O_2S$, the The C=S group lies on a mirror plane. The molecules are packed in a centrosymmetric manner through intermolecular $N-H\cdots$ S hydrogen bonds.

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Comment

Compounds containing the thiourea unit show strong antifungal and antibacterial activity and find applications in medicine and agriculture (Lin *et al.*, 2004). The title compound, (I), crystallized in the space group *Pnma* with Z =4. The C=S group lies on a mirror plane and thus there is one half-molecule in the asymmetric unit.



The molecule of (I) is shown in Fig. 1 and a packing diagram is shown in Fig. 2. The dihedral angle between the planes formed by N1/C1/S1 and C2–C7 is 90.26 (1)°, resulting in a bent 'butterfly-like' molecular shape. The molecules are arranged parallel to each other and extend along the long *b* axis *via* purely due to purely van der Waals interactions. Two well defined N–H···S hydrogen bonds involving the S atom lying on the mirror plane generate the packing motif in the *ac* plane [N–H···S: N···S = 3.551 (3) Å]. As a result, alternate molecules generated by the glide planes are held perpendicular to each other, as shown in Fig. 2.

Experimental

To a solution of *p*-anisidine (37 g), carbon disulfide (24 ml) and rectified spirit (40 ml) kept at 283–287 K, a small amount of aqueous ammonia (41 ml) was added, accompained by constant shaking. The intermediate thiocarbamate was recovered and washed with small amounts of diethyl ether. It was subjected to steam distillation after adding lead nitrate and water. The compound which separated was further distilled and crystallized. The NMR data for the compound have been reported previously (Natarajan *et al.*, 2005).

Crystal data $C_{15}H_{16}N_2O_2S$ $M_r = 288.37$ Orthorhombic, Pnma a = 8.427 (5) Å b = 31.628 (19) Å c = 5.292 (3) Å V = 1410.3 (14) Å³ Z = 4 $D_r = 1.358$ Mg m⁻³

Mo K α radiation Cell parameters from 732 reflections $\theta = 2.6-27.4^{\circ}$ $\mu = 0.23$ mm⁻¹ T = 290 (2) K Prism, colourless 0.45 × 0.40 × 0.35 mm

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Figure 1

View of the molecular structure of (I), showing the atom-labelling scheme. Displacement ellipsoids are drawn at the 50% probability level. H atoms are shown as small spheres of arbitrary radii. [Symmetry code: (a) $x, \frac{1}{2} - y, z.$]

Data collection

Bruker SMART CCD area-detector	1572 independent reflections
diffractometer	1433 reflections with $I > 2\sigma(I)$
φ and ω scans	$R_{\rm int} = 0.022$
Absorption correction: multi-scan	$\theta_{\rm max} = 27.4^{\circ}$
(SADABS; Sheldrick, 1996)	$h = -10 \rightarrow 10$
$T_{\min} = 0.903, T_{\max} = 0.923$	$k = -40 \rightarrow 39$
11344 measured reflections	$l = -6 \rightarrow 6$

Refinement

$w = 1/[\sigma^2(F_o^2) + (0.0502P)^2]$
+ 0.7475P]
where $P = (F_0^2 + 2F_c^2)/3$
$(\Delta/\sigma)_{\rm max} < 0.001$
$\Delta \rho_{\rm max} = 0.33 \ {\rm e} \ {\rm \AA}^{-3}$
$\Delta \rho_{\rm min} = -0.19 \text{ e} \text{ \AA}^{-3}$

Table 1

Selected geometric parameters (Å, °).

S1-C1	1.693 (3)	N1-C1	1.339 (2)
O1-C5	1.375 (3)	N1-C2	1.431 (3)
O1-C8	1.417 (4)		
C5-O1-C8	117.5 (2)	N1-C2-C3	119.41 (19)
C1-N1-C2	124.68 (18)	N1-C2-C7	120.56 (18)
N1-C1-N1 ⁱ	114.7 (2)	O1-C5-C6	124.4 (2)
S1-C1-N1	122.66 (12)	O1-C5-C4	115.5 (2)
-	. ,		

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

Table 2

Hydrogen-bond geometry (Å, °).

$D - H \cdot \cdot \cdot A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdot \cdot \cdot A$
$N1-H1\cdots S1^{ii}$	0.80 (3)	2.79 (3)	3.551 (3)	160 (2)
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Symmetry code: (ii) $x + \frac{1}{2}, -y + \frac{1}{2}, -z + \frac{1}{2}$.

All the H atoms were located and refined isotropically. The C-H bond lengths are 0.89(3)-1.05(3) Å and the N-H bond length is 0.80 (3) Å.

Data collection: SMART (Bruker, 1998); cell refinement: SMART; data reduction: SAINT (Bruker, 1998); program(s) used to solve structure: SHELXS97 (Sheldrick, 1997); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics:



Figure 2

Packing diagram of (I), viewed down the c axis. The dotted lines indicate intermolecular N-H···S hydrogen bonds.

ORTEP-3 for Windows (Farrugia, 1997) and CAMERON (Watkin et al., 1993); software used to prepare material for publication: PLATON (Spek, 2003).

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