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Key indicators

Single-crystal X-ray study T = 293 K Mean σ (C–C) = 0.003 Å R factor = 0.036 wR factor = 0.101 Data-to-parameter ratio = 14.6

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

{3-[4-(*N*,*N*-Dimethylamino)phenyl]-1-(4-methyl-phenyl)-3-(phenylsulfanyl)propylidene}semicarbazide

In the title compound, $C_{25}H_{28}N_4OS$, the benzene rings are nearly perpendicular to the central C-C-C-S linkage, whereas the semicarbazide plane forms a dihedral angle of 67.7 (1)° with the central linkage. In the crystal structure, inversion-related molecules exist as N-H···O hydrogenbonded dimers. Received 27 August 2004 Accepted 23 September 2004 Online 30 September 2004

Comment

Studies on 1,2,3-selenadiazoles and 1,2,3-thiadiazoles have attracted much attention not only due to their pharmacological properties, such as antifungal (Jalilian et al., 2003) and antibacterial activities (Lalezari et al., 1974), but also due to their wide-ranging utility as synthons in organic synthesis (Arsenyan et al., 2002; Morzherin et al., 2003). In a continuation of our work on the synthesis and characterization of 1,2,3selenadiazoles and 1,2,3-thiadiazoles (Muthusubramanian et al., 1997; Saravanan & Muthusubramanian, 2004), we planned to synthesize more 1,2,3-selena/thiadiazoles. In this regard, we prepared several semicarbazones of 1,3-diaryl-3-phenylthiopropan-1-ones, including the title compound, $\{3-[4-(N,N-di$ methylamino)phenyl]-1-(4-methylphenyl)-3-(phenylsulfanyl)propylidene}semicarbazide, (I). The reaction of these ketones with semicarbazide hydrochloride is very sensitive in the sense that heating the reaction mixture leads to elimination of the phenylsulfanyl group, and hence the reaction was carried out at room temperature in dioxane in the presence of the phasetransfer catalyst tetrabutylammonium bromide.



In the title molecule, the phenylsulfanyl (C20–C25), methylphenyl (C2–C7), dimethylaminophenyl (C12–C17) and semicarbazide (N1–N3/O1/C9) groups make dihedral angles of 87.9 (1), 88.2 (1), 83.0 (1) and 67.7 (1)°, respectively, with the central C8–C10–C11–S1 linkage. The C8–C10–C11–C12 and C8–C10–C11–S1 torsion angles of 72.3 (2) and -165.8 (1)°, respectively, show that the imino C atom is nearly *gauche* to the C12-aryl ring and *anti* to the phenylsulfanyl ring.

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Figure 1

A view of the molecule of of (I), showing the atom-numbering scheme. Displacement ellipsoids are drawn at the 50% probability level.



Figure 2

Packing of the molecules, viewed down the *a* axis, showing the dimer formation. Hydrogen bonds are indicated by dashed lines.

The close approach of atoms H2 and H10B (H2···H10B = 1.97 Å) results in the widening and narrowing of angles N1- $C8-10 [127.6 (2)^{\circ}]$ and $N1-C8-C5 [113.9 (2)^{\circ}]$ from the normal 120°.

In the crystalline state, inversion-related molecules exist as $N2-H2\cdotsO1^{i}$ and $C10-H10B\cdotsO1^{i}$ hydrogen-bonded dimers (Table 1). One of the amino H atoms, H3B, is involved in an intramolecular N3-H3B···N1 hydrogen bond, whereas the other atom, H3A, is not involved in any hydrogen bond (Fig. 2).

Experimental

To a solution of 3-[4-(dimethylamino)phenyl]-1-(4-methylphenyl)-3-(phenylsulfanyl)propan-1-one (1.88 g, 0.005 mol), semicarbazide hydrochloride (4.46 g) and sodium acetate (3.28 g) in a dioxanewater mixture (50 ml, 3:2 v/v), a catalytic amount of the phasetransfer catalyst tetrabutylammonium bromide was added. The reaction mixture was stirred at room temperature for 3 d, then extracted with chloroform and evaporated to dryness. The insoluble ethyl acetate portion was crystallized from ethanol to give the title compound (m.p. 445 K). IR (KBr, v, cm⁻¹): 3508, 3392, 1682, 1562; ¹H NMR (CDCl₃, p.p.m.): 9.33 (s, 1H), 8.13 (s, 2H), 7.39 (d, 2H, J = 8.1 Hz), 7.12–7.19 (m, 5H), 6.56 (d, 2H, J = 8.7 Hz), 6.29 (s, 2H), 4.33 (t, 1H), 3.65 (m, 2H), 2.86 (s, 6H), 2.33 (s, 3H); ¹³C NMR (CDCl₃, p.p.m.): 157.70, 149.89, 145.40, 138.17, 135.15, 134.62, 131.71, 129.04, 128.98, 128.40, 128.10, 127.17, 126.45, 112.20, 79.14, 50.18, 33.21, 21.15.

Crystal data

$C_{25}H_{28}N_4OS$	Z = 2
$M_r = 432.57$	$D_x = 1.233 \text{ Mg m}^{-3}$
Triclinic, P1	Mo $K\alpha$ radiation
a = 6.804 (5) Å	Cell parameters from 25
b = 10.531 (4) Å	reflections
c = 17.278 (5) Å	$\theta = 11.2 - 14.1^{\circ}$
$\alpha = 88.96 \ (5)^{\circ}$	$\mu = 0.16 \text{ mm}^{-1}$
$\beta = 79.24 \ (3)^{\circ}$	T = 293 (2) K
$\gamma = 73.51 \ (2)^{\circ}$	Needle, colourless
$V = 1165.4 (10) \text{ Å}^3$	$0.35 \times 0.25 \times 0.15 \text{ mm}$

 $R_{\rm int} = 0.014$

 $\theta_{\rm max} = 25.0^\circ$

 $h = -7 \rightarrow 8$

 $k = -1 \rightarrow 12$

 $l = -20 \rightarrow 20$

3 standard reflections

frequency: 60 min

intensity decay: none

Data collection

Nonius MACH3 four-circle diffractometer ω -2 θ scans Absorption correction: ψ scan (North et al., 1968) $T_{\rm min}=0.975,\ T_{\rm max}=0.982$ 4584 measured reflections 4083 independent reflections 3120 reflections with $I > 2\sigma(I)$

Refinement

Refinement on F^2	$w = 1/[\sigma^2(F_o^2) + (0.0454P)^2]$
$R[F^2 > 2\sigma(F^2)] = 0.036$	+ 0.3387P]
$wR(F^2) = 0.101$	where $P = (F_o^2 + 2F_c^2)/3$
S = 1.03	$(\Delta/\sigma)_{\rm max} < 0.001$
4083 reflections	$\Delta \rho_{\rm max} = 0.19 \ {\rm e} \ {\rm \AA}^{-3}$
280 parameters	$\Delta \rho_{\rm min} = -0.22 \text{ e } \text{\AA}^{-3}$
H-atom parameters constrained	

Table 1 Hydrogen-bonding geometry (Å, °).

$D - H \cdots A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - H \cdots A$
$N2 - H2 \cdots O1^{i}$ $C10 - H10B \cdots O1^{i}$ $N3 - H3B \cdots N1$	0.86	2.03	2.843 (3)	157
	0.97	2.38	3.333 (2)	169
	0.86	2.24	2.606 (2)	105

Symmetry code: (i) 2 - x, 1 - y, 1 - z.

All H atoms were placed in calculated positions (N-H = 0.86 Å and C-H = 0.93-0.98 Å) and included in the refinement in a ridingmodel approximation, with U_{iso} values constrained to be $1.5U_{eq}$ of the carrier atom for methyl H atoms and $1.2U_{eq}$ for the remaining H atoms.

Data collection: *CAD-4 EXPRESS* (Enraf–Nonius, 1994); cell refinement: *CAD-4 EXPRESS*; data reduction: *XCAD*4 (Harms & Wocadlo, 1995); program(s) used to solve structure: *SHELXS97* (Sheldrick, 1990); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *PLATON* (Spek, 1999); software used to prepare material for publication: *SHELXL97*.

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