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Key indicators

Single-crystal X-ray study T = 293 K Mean σ (C–C) = 0.002 Å R factor = 0.044 wR factor = 0.126 Data-to-parameter ratio = 19.9

For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e. The title molecule, $C_9H_{11}N_3OS$, exhibits a *trans* conformation with respect to the phenyl ring and the thiosemicarbazone moiety. The dihedral angle between the thiosemicarbazone moiety and the phenyl ring is 4.68 (5)°. In the solid state, inversion-related molecules exist as centrosymmetric N-H···S hydrogen-bonded dimers. Symmetry-related dimers are interlinked by N-H···S and N-H···O intermolecular hydrogen bonds, forming two-dimensional molecular networks parallel to the *ab* plane.

2-Methoxybenzaldehyde thiosemicarbazone

Comment

Thiosemicarbazones are derivatives of carbonyl compounds, exhibiting a wide range of biological activities (Shanmuga Sundara Raj *et al.*, 2000). There is considerable interest in the chemistry of Schiff base compounds containing N- and Sdonors and of their metal complexes. This is due to their nonlinear optical properties and chelating ability with transition metal ions (Ali & Tarafdar, 1977; Ali & Bose, 1984; Tian *et al.*, 1996). Antibacterial, antiviral and antitumour activities have been observed in N- and S-donor ligands, such as substituted thiosemicarbazides and thiosemicarbazones (Nandi *et al.*, 1984; French & Blanz, 1965, 1966; William, 1972). In this paper, we report the structure of 2-methoxybenzaldehyde thiosemicarbazone, (I).



The molecular structure of (I), with the atom-numbering scheme, is shown in Fig. 1. The bond distances agree well with the values reported for similar structures (Moers et al., 1999; Nandi et al., 1984; Shanmuga Sundara Raj et al., 2000). The C=S bond [1.688 (2) Å] has a length intermediate between a single and double bond. The sum of the valence angles around atoms N1 and N2 indicate that these atoms are sp^2 hybridized. The exocyclic angles around atom C4 show considerable asymmetry, with the O1-C4-C5 angle $[123.9 (2)^{\circ}]$ being wider than O1-C4-C3 [115.9 (1) $^{\circ}$], as observed in a related structure (Fun et al., 1996). This may be due to the steric repulsion between the methyl group and the phenyl ring. The C9-O1-C4-C5 [-14.1 (3)°] and C9-O1-C4-C3 $[166.4 (2)^{\circ}]$ torsion angles indicate that the methoxy group is twisted away from the benzene plane. The S1-C1-N2-N3 $[-177.6 (1)^{\circ}]$ and N3-C2-C3-C4 $[-178.1 (2)^{\circ}]$ torsion

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Figure 1 The molecular structure of (I), with the atom-numbering scheme. Displacement ellipsoids are shown at the 50% probability level.

angles indicate a trans conformation with respect to the thiosemicarbazone moiety and the phenyl ring. The trans conformation adopted by the side chain is evident from the values of the C1-N2-N3-C2 [172.8 (2)°], N2-N3-C2-C3 $[-178.6 (1)^{\circ}]$, N3-C2-C3-C4 $[-178.1 (2)^{\circ}]$ and C2-C3-C4-C5 $[-178.4 (2)^{\circ}]$ torsion angles. The thiosemicarbazone moiety forms a dihedral angle of $4.68 (5)^{\circ}$ with the phenyl ring.

In the crystal, the molecules at (x, y, z) and (-x, 1-y, z)(1 - z) are linked by N-H···S hydrogen bonds involving the NH group, forming centrosymmetric dimeric units. Symmetryrelated dimers are interlinked by $N-H\cdots S$ and $N-H\cdots O$ intermolecular hydrogen bonds involving the NH₂ group (Table 2), forming two-dimensional molecular networks parallel to the *ab* plane (Fig. 2). The $N \cdots S$ distances of 3.376 (1) and 3.398 (1) Å are close to the mean $N \cdot \cdot S$ distance reported for N-H···S hydrogen bonds by Srinivasan & Chacko (1967).

Experimental

2-Methoxybenzaldehyde and thiosemicarbazide, in almost equivalent quantities, with a few drops of HCl, yielded 2-methoxybenzaldehyde thiosemicarbazone. Single crystals suitable for X-ray diffraction analysis were obtained by slow evaporation from an acetone solution at room temperature.

Crystal data

C ₉ H ₁₁ N ₃ OS	Mo $K\alpha$ radiation
$M_r = 209.27$	Cell parameters from 6556
Orthorhombic, Pbca	reflections
a = 13.5826 (2) Å	$\theta = 2.8-28.3^{\circ}$
b = 10.6063 (2) Å	$\mu = 0.28 \text{ mm}^{-1}$
c = 14.3260 (3) Å	T = 293 (2) K
V = 2063.82 (7) Å ³	Block, colourless
Z = 8	$0.48 \times 0.34 \times 0.20 \text{ mm}$
$D_x = 1.347 \text{ Mg m}^{-3}$	
Data collection	
Siemens SMART CCD area-	1742 reflections with $I > 2\sigma(I)$
detector diffractometer	$R_{\rm int} = 0.079$
ω scans	$\theta_{\rm max} = 28.3^{\circ}$
Absorption correction: none	$h = -18 \rightarrow 17$
13626 measured reflections	$k = -14 \rightarrow 10$
2547 independent reflections	$l = -19 \rightarrow 13$



Figure 2

The packing of the title molecules, viewed down the b axis.

Refinement

Refinement on F^2	$w = 1/[\sigma^2(F_o^2) + (0.0674P)^2]$
$R[F^2 > 2\sigma(F^2)] = 0.044$	where $P = (F_o^2 + 2F_c^2)/3$
$wR(F^2) = 0.126$	$(\Delta/\sigma)_{\rm max} < 0.001$
S = 0.98	$\Delta \rho_{\rm max} = 0.32 \ {\rm e} \ {\rm \AA}^{-3}$
2547 reflections	$\Delta \rho_{\rm min} = -0.34 \text{ e } \text{\AA}^{-3}$
128 parameters	Extinction correction: SHELXL97
H-atom parameters constrained	Extinction coefficient: 0.0093 (15)

Table 1

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Selected geometric parameters (Å, °).

S1-C1	1.688 (2)	N2-C1	1.345 (2)
O1-C4	1.377 (2)	N2-N3	1.377 (2)
O1-C9	1.445 (2)	N3-C2	1.276 (2)
N1-C1	1.321 (2)	C2-C3	1.460 (2)
O1-C4-C5	123.9 (2)	01-C4-C3	115.9 (1)
C1-N2-N3-C2	172.8 (2)	C9-O1-C4-C5	-14.1 (3)
N3-N2-C1-S1	-177.6(1)	C9-O1-C4-C3	166.4 (2)
N2-N3-C2-C3	-178.6(1)	C2-C3-C4-C5	-178.4(2)
N3-C2-C3-C4	-178.1(2)		

Table 2	
Hydrogen-bonding geometry	(Å, °).

$D - H \cdots A$	$D-\mathrm{H}$	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdots A$
$C2-H2A\cdots O1$	0.93	2.42	2.7427 (19)	100
$N1 - H1A \cdot \cdot \cdot N3$	0.86	2.29	2.6385 (19)	105
$N1 - H1B \cdot \cdot \cdot O1^{i}$	0.86	2.19	3.0306 (18)	166
$N1-H1A\cdots S1^{ii}$	0.86	2.73	3.3982 (14)	136
$N2-H2\cdots S1^{iii}$	0.86	2.58	3.3759 (14)	154

Symmetry codes: (i) $\frac{1}{2} + x, \frac{3}{2} - y, 1 - z$; (ii) $\frac{1}{2} - x, \frac{1}{2} + y, z$; (iii) -x, 1 - y, 1 - z.

All H atoms were positioned geometrically and were treated as riding on their parent C and N atoms, with C-H distances of 0.93 or 0.96 Å and N-H distances of 0.86 Å.

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Data collection: *SMART* (Siemens, 1996); cell refinement: *SAINT* (Siemens, 1996); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS*97 (Sheldrick, 1997); program(s) used to refine structure: *SHELXL*97 (Sheldrick, 1997); molecular graphics: *ORTEP*-3 (Farrugia, 1997) and *PLATON* (Spek, 1990); software used to prepare material for publication: *SHELXL*97 and *PARST* (Nardelli, 1995).

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