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## Key indicators

Single-crystal X-ray study T = 173 K Mean  $\sigma$ (C–C) = 0.002 Å R factor = 0.031 wR factor = 0.084 Data-to-parameter ratio = 22.9

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

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© 2006 International Union of Crystallography All rights reserved The title compound,  $C_9H_{11}N_3S_2$ , crystallizes with two molecules in the asymmetric unit, which differ mainly in the degree of planarity. The C=N double bonds are *trans* configured. Geometric parameters are in the usual ranges. The crystal structure is characterized by N-H···N and N-H···S hydrogen bonds.

4-(Methylsulfanyl)benzaldehyde thiosemicarbazone

# Comment

Thiosemicarbazones (TSCs) of aromatic aldehydes and ketones are widely known as carcinostatic and antimicrobial agents. It has been found that silyl-substituted furfural TSCs possess neurotropic activity (Lukevlcs *et al.*, 1993). TSCs of arylidene and aryl aldehydes and ketones have shown anticonvulsant activity in maximal electroshock seizure tests (Dimmock *et al.*, 1986, 1990, 1991). In view of the importance of thiosemicarbazones, a new thiosemicarbazone, (I), has been prepared and its crystal structure is reported here.



A perspective view of (I) is shown in Fig. 1. Bond lengths and angles can be regarded as normal (Cambridge Structural Database, Version 5.27, November 2005 updated August 2006; MOGUL Version 1.1; Allen, 2002). The C=N double bonds are trans configured. Whereas one molecule (labelled with suffix A) is essentially planar (r.m.s. deviation = 0.062 Å for all non-H atoms) the other one deviates markedly from planarity (r.m.s. deviation = 0.262 Å for all non-H atoms). The methylsulfanyl residues are coplanar with the benzene rings (Table 1). The thiosemicarbazone units are planar (r.m.s. deviation = 0.036 Å for the non-planar molecule and 0.014 Å for molecule A), but they form different dihedral angles with the benzene ring, viz. 25.19 (4)° for the non-planar molecule and 7.32 (6)° for molecule A. Both molecules show an intramolecular N-H···N contact and two N-H···S hydrogen bonds (Table 2). It is remarkable that only the thiocarbonyl S atom acts as an acceptor for hydrogen bonds, while the methylsulfanyl S atom is not involved in hydrogen bonds.

# **Experimental**

A mixture of 4-methylsulfanylbenzaldehyde (1.52 g, 0.01 mol) and thiosemicarbazide (0.91 g, 0.01 mol) in ethanol (15 ml) was refluxed

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## Figure 1

Perspective view of the two molecules in the asymmetric unit of the title compound with the atom numbering; displacement ellipsoids are drawn at the 50% probability level.

for 3 h on a water bath. The precipitated solid was filtered, washed with water, dried and recrystallized from acetone (yield: 81%; m.p. 463-465 K). Analysis (%) found (calculated) for C<sub>9</sub>H<sub>11</sub>N<sub>3</sub>S<sub>2</sub>: C 47.84 (47.97), H 4.85 (4.92), N 18.54 (18.65), S 28.38 (28.46).

## Crystal data

$C_9H_{11}N_3S_2$	$V = 1089.76 (10) \text{ Å}^3$
$M_r = 225.33$	Z = 4
Triclinic, $P\overline{1}$	$D_x = 1.373 \text{ Mg m}^{-3}$
a = 8.2052 (4) Å	Mo $K\alpha$ radiation
b = 9.1947 (5)  Å	$\mu = 0.45 \text{ mm}^{-1}$
c = 15.5994 (8) Å	T = 173 (2) K
$\alpha = 106.558 \ (5)^{\circ}$	Block, light orange
$\beta = 97.573 \ (3)^{\circ}$	$0.38 \times 0.36 \times 0.33$ r
$\gamma = 100.110 \ (4)^{\circ}$	

#### Data collection

Stoe IPDS-II two-circle diffractometer  $\omega$  scans Absorption correction: multi-scan (MULABS; Spek, 2003; Blessing, 1995)  $T_{\rm min}=0.847,\ T_{\rm max}=0.865$ 

#### Refinement

Refinement on  $F^2$  $R[F^2 > 2\sigma(F^2)] = 0.031$ wR(F<sup>2</sup>) = 0.084 S = 1.066421 reflections 280 parameters H atoms treated by a mixture of independent and constrained refinement

 $Mg m^{-3}$ diation  $nm^{-1}$ 2) K nt orange  $6 \times 0.33$  mm

36432 measured reflections 6421 independent reflections 6026 reflections with  $I > 2\sigma(I)$  $R_{\rm int} = 0.041$  $\theta_{\rm max} = 30.2^\circ$ 

 $w = 1/[\sigma^2(F_0^2) + (0.0409P)^2]$ + 0.3428P] where  $P = (F_o^2 + 2F_c^2)/3$  $(\Delta/\sigma)_{\rm max} = 0.001$  $\Delta \rho_{\rm max} = 0.38 \text{ e } \text{\AA}^{-3}$  $\Delta \rho_{\rm min} = -0.28 \text{ e } \text{\AA}^{-3}$ Extinction correction: SHELXL97 Extinction coefficient: 0.048 (3)

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Selected torsion angles (°).

C3-S2-C14-C15	5.08 (12)	C3A-S2A-C14A-C15A	7.41 (12)

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

$D - H \cdot \cdot \cdot A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdots A$
$N1-H1A\cdots S1^{i}$	0.873 (17)	2.459 (17)	3.3254 (11)	172.1 (15)
$N1 - H1B \cdot \cdot \cdot N3$	0.875 (17)	2.266 (16)	2.6364 (13)	105.4 (12)
$N2-H2\cdots S1A^{i}$	0.854 (17)	2.555 (18)	3.3412 (10)	153.5 (15)
$N1A - H1C \cdot \cdot \cdot S1A^{ii}$	0.869 (19)	2.604 (19)	3.4707 (11)	174.7 (17)
$N2A - H2A \cdot \cdot \cdot S1^{i}$	0.906 (17)	2.454 (17)	3.3499 (10)	169.8 (14)
$N1A - H1D \cdots N3A$	0.852 (17)	2.302 (16)	2.6440 (14)	104.2 (13)

Symmetry codes: (i) -x + 1, -y, -z + 1; (ii) -x + 1, -y, -z.

H atoms were found in a difference map. Those bonded to carbon were refined using a riding model, with C-H = 0.95 Å for aromatic and C-H = 0.98 Å for methyl groups.  $U_{iso}(H)$  values were set at  $1.2U_{eq}(C)$  [1.5 $U_{eq}(methyl C)$ ]. In addition, the methyl groups were allowed to rotate but not to tip. H atoms bonded to N were freely refined.

Data collection: X-AREA (Stoe & Cie, 2001); cell refinement: X-AREA; data reduction: X-AREA; program(s) used to solve structure: SHELXS97 (Sheldrick, 1997); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: PLATON (Spek, 2003); software used to prepare material for publication: SHELXL97 and PLATON.

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