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## 4-(Methylsulfanyl)benzaldehyde thiosemicarbazone

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#### **Key indicators**

Single-crystal X-ray study  $T=173~\mathrm{K}$  Mean  $\sigma(\mathrm{C-C})=0.002~\mathrm{\mathring{A}}$  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.

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  $N-H\cdots N$  and  $N-H\cdots S$  hydrogen bonds.

### 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  $N{-}H{\cdot}{\cdot}{\cdot}N$  contact and two  $N{-}H{\cdot}{\cdot}{\cdot}S$  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**

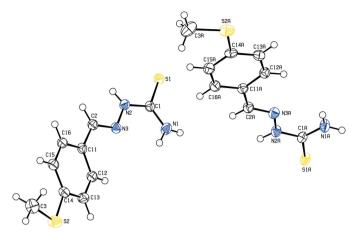
A mixture of 4-methylsulfanylbenzaldehyde (1.52 g, 0.01 mol) and thiosemicarbazide (0.91 g, 0.01 mol) in ethanol (15 ml) was refluxed

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## organic papers



**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  $C_9H_{11}N_3S_2$ : C 47.84 (47.97), H 4.85 (4.92), N 18.54 (18.65), S 28.38 (28.46).

## Crystal data

$V = 1089.76 (10) \text{ Å}^3$
Z = 4
$D_x = 1.373 \text{ Mg m}^{-3}$
Mo $K\alpha$ radiation
$\mu = 0.45 \text{ mm}^{-1}$
T = 173 (2)  K
Block, light orange
$0.38 \times 0.36 \times 0.33 \text{ mm}$

## Data collection

Stoe IPDS-II two-circle diffractometer diffractometer 6421 independent reflections  $\omega$  scans 6026 reflections with  $I > 2\sigma(I)$  Absorption correction: multi-scan (MULABS; Spek, 2003; Blessing, 1995)  $T_{\min} = 0.847, T_{\max} = 0.865$ 

### Refinement

Refinement on $F^2$	$w = 1/[\sigma^2(F_0^2) + (0.0409P)^2]$
$R[F^2 > 2\sigma(F^2)] = 0.031$	+ 0.3428P]
$wR(F^2) = 0.084$	where $P = (F_0^2 + 2F_c^2)/3$
S = 1.06	$(\Delta/\sigma)_{\text{max}} = 0.001$
6421 reflections	$\Delta \rho_{\text{max}} = 0.38 \text{ e Å}^{-3}$
280 parameters	$\Delta \rho_{\min} = -0.28 \text{ e Å}^{-3}$
H atoms treated by a mixture of	Extinction correction: SHELXL97
independent and constrained	Extinction coefficient: 0.048 (3)
refinement	

**Table 1** Selected torsion angles (°).

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

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

$D-H\cdots A$	D-H	$H \cdot \cdot \cdot A$	$D \cdot \cdot \cdot A$	$D-\mathrm{H}\cdots A$
$N1-H1A\cdots S1^{i}$	0.873 (17)	2.459 (17)	3.3254 (11)	172.1 (15)
$N1-H1B\cdots N3$	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)
$N1A - H1D \cdot \cdot \cdot N3A$	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.  $U_{\rm iso}({\rm H})$  values were set at  $1.2 U_{\rm eq}({\rm C})$  [1.5 $U_{\rm eq}({\rm methyl}$  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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