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#### **Key indicators**

Single-crystal X-ray study T = 293 K Mean  $\sigma$ (C–C) = 0.003 Å R factor = 0.041 wR factor = 0.103 Data-to-parameter ratio = 15.2

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

# Bis(4-formyl-3-methyl-1-phenyl-1*H*-pyrazol-5-yl) sulfide

The molecule of the title compound,  $C_{22}H_{18}N_4O_2S$ , is nonplanar and possesses no crystallographic symmetry. The crystal structure involves intermolecular  $C-H\cdots O$  hydrogen bonds. Received 26 August 2004 Accepted 24 September 2004 Online 30 September 2004

### Comment

Pyrazole sulfide derivatives exhibit good fungicidal activity (Li *et al.*, 1997; Yu *et al.*, 1999). In an effort to obtain lead compounds, we have synthesized a series of such derivatives.



The crystal structure of (I) has been determined and the results are presented here. The molecular structure is shown in Fig. 1. The dihedral angle between the N1-pyrazole ring plane and its attached phenyl ring is 78.41 (7)°; that between the N3-pyrazole ring and its attached phenyl ring is  $61.90 (9)^{\circ}$ . The dihedral angle between the two pyrazole rings is  $75.10 (8)^{\circ}$ . Selected bond lengths and angles are listed in Table 1. These are in excellent agreement with the numerous literature reports of pyrazole derivatives (Allen *et al.*, 1987). The crystal structure involves intermolecular C–H···O hydrogen bonds (Table 2).



# hy The mole

© 2004 International Union of Crystallography Printed in Great Britain – all rights reserved The molecular structure of (I), showing the atom-labellling scheme. Displacement ellipsoids are drawn at the 50% probability level.



#### Figure 2

The molecular packing of (I), viewed along the *b* axis. Dashed lines indicate hydrogen bonds.

## **Experimental**

To a 100 ml nitrogen-flushed three-neck round-bottomed flask, the catalyst NaH (10 mmol) in DMF was added. Thiourea (10 mmol) in DMF (10 ml) was added dropwise. After 30 min, 5-chloro-3-methyl-1-phenyl-pyrazol-4-yl carboxaldehyde was added dropwise and the solution was stirred for 24 h, under N2 at 383 K. The reaction gave a mixture containing (I), which was separated by recrystallization from ethanol. Crystals suitable for X-ray analysis were obtained by evaporation of an ethanol solution (m.p. 468-469 K). <sup>1</sup>H NMR (DMSO, 400 MHz): δ 9.48 (s, 2H, CH=O), 7.31-7.43 (m, 10H, C<sub>6</sub>H<sub>5</sub>, C<sub>6</sub>H<sub>3</sub>), 2.19 (s, 6H, CH<sub>3</sub>); L-MS (IS, 4500; TEM, 0.00; GS1, 14.00; GS2, 0.00) (m/z, relative intensity %): 425 (M + Na, 100.00), 403 (M + 1, 70.07), 402 (M, 1.75), 213 (1.94). Analysis calculated for C22H18N4O2S: C 65.60, H 4.47, N 13.91%; found: C 65.78, H 4.61, N 14.12%.

## Crystal data C22H18N4O2S

$M_r = 402.46$
Monoclinic, $P2_1/c$
a = 13.853 (4)  Å
b = 6.918 (2)  Å
c = 20.723 (6) Å
$\beta = 99.855 \ (5)^{\circ}$
$V = 1956.6 (10) \text{ Å}^3$
Z = 4
Data collection
Siemens SMART CCD area-
detector diffractometer
$\varphi$ and $\omega$ scans
Absorption correction: multi-scan
(SADABS; Sheldrick, 1996)
$T_{\rm min} = 0.918, T_{\rm max} = 0.933$

10 858 measured reflections

 $D_x = 1.366 \text{ Mg m}^{-3}$ Mo  $K\alpha$  radiation Cell parameters from 894 reflections  $\theta = 3.4 - 24.6^{\circ}$  $\mu = 0.19~\mathrm{mm}^{-1}$ T = 293 (2) KBlock, colorless  $0.42 \times 0.40 \times 0.36 \text{ mm}$ 

#### 4015 independent reflections 2913 reflections with $I > 2\sigma(I)$ $R_{\rm int}=0.029$ $\theta_{\rm max} = 26.5^{\circ}$ $h = -17 \rightarrow 17$ $k = -4 \rightarrow 8$ $l = -25 \rightarrow 22$

Refinement

T.L.L. 4

Refinement on $F^2$	$w = 1/[\sigma^2(F_o^2) + (0.045P)^2]$
$R[F^2 > 2\sigma(F^2)] = 0.041$	+ 0.5946P]
$wR(F^2) = 0.103$	where $P = (F_o^2 + 2F_c^2)/3$
S = 1.02	$(\Delta/\sigma)_{\rm max} = 0.001$
4015 reflections	$\Delta \rho_{\rm max} = 0.21 \ {\rm e} \ {\rm \AA}^{-3}$
265 parameters	$\Delta \rho_{\rm min} = -0.19 \text{ e } \text{\AA}^{-3}$
H-atom parameters constrained	Extinction correction: SHELXL97
	Extinction coefficient: 0.0124 (10)

Table T				
Selected	geometric	parameters	(Å,	°).

-			
S1-C14	1.7555 (19)	N4-C14	1.355 (2)
S1-C3	1.7556 (18)	O1-C5	1.214 (2)
N1-C1	1.329 (2)	O2-C16	1.205 (2)
N1-N2	1.3724 (19)	C1-C2	1.416 (3)
N2-C3	1.344 (2)	C2-C3	1.384 (2)
N3-C12	1.324 (2)	C12-C13	1.411 (3)
N3-N4	1.374 (2)	C13-C14	1.382 (2)
C14 - S1 - C3	99.24 (8)	$C_{3}-C_{2}-C_{1}$	104.63 (16)
C1-N1-N2	105.00 (15)	N2-C3-C2	107.31 (15)
C3-N2-N1	111.78 (14)	N3-C12-C13	111.64 (16)
C12-N3-N4	104.76 (16)	C14-C13-C12	105.03 (17)
C14-N4-N3	111.85 (15)	N4-C14-C13	106.68 (17)
N1-C1-C2	111.26 (15)		

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

$D - H \cdots A$	$D-\mathrm{H}$	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdots A$
$C9 - H9 \cdots O1^{i}$	0.93	2.49	3.244 (3)	138
C11−H11···O1 <sup>ii</sup>	0.93	2.57	3.417 (3)	151
$C21 - H21 \cdots O2^{iii}$	0.93	2.53	3.385 (3)	153

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

H atoms were placed in calculated positions, with C-H distances of 0.93 and 0.96 Å.  $U_{iso}(H)$  values were constrained to  $1.5U_{eq}(carrier)$ atom) for methyl H atoms and  $1.2U_{eq}$ (carrier atom) for other H atoms.

Data collection: SMART (Bruker, 1997-2002); cell refinement: SMART; data reduction: SAINT (Bruker, 1997-2002); program(s) used to solve structure: SHELXS97 (Sheldrick, 1997); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: SHELXTL (Bruker, 1997-2002); software used to prepare material for publication: SHELXTL.

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