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## Key indicators

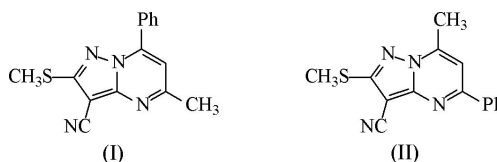
Single-crystal X-ray study  
 $T = 293$  K  
Mean  $\sigma(\text{C}-\text{C}) = 0.004$  Å  
 $R$  factor = 0.051  
 $wR$  factor = 0.122  
Data-to-parameter ratio = 15.7For details of how these key indicators were  
automatically derived from the article, see  
<http://journals.iucr.org/e>.5-Methyl-2-methylsulfanyl-7-phenylpyrazolo-  
[1,5-*a*]pyrimidine-3-carbonitrileThe pyrazolo[1,5-*a*]pyrimidine ring system of the title  
compound,  $\text{C}_{15}\text{H}_{12}\text{N}_4\text{S}$ , is essentially planar and the dihedral  
angle between the plane of the pyrazolo[1,5-*a*]pyrimidine  
system and the plane of the phenyl ring is  $134.6$  ( $7$ )°.

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## Comment

Pyrazolo[1,5-*a*]pyrimidines are of pharmacological impor-  
tance as purine analogues and, used in the treatment of  
hyperuricaemia and gout, inhibit *de novo* purine biosynthesis  
and xanthine oxidase (El-Gaby *et al.*, 2000; Elnagdi *et al.*,  
1987). In our recent research, a series of derivatives have been  
synthesized in order to look for new compounds having  
herbicidal activity (Sawhney *et al.*, 1981). Cyclization of  
5-aminopyrazole with a dione compound gives the desired  
pyrazolo[1,5-*a*]pyrimidine, which results in the production of  
two positional isomers, (I) and (II).The structure of (I), reported here (Fig. 1), allowed us to  
investigate the relationship between structure and herbicidal  
activity. There are three planes in (I); these are defined as  $p1$   
(C1/C2/C3/N1/N2),  $p2$  (C3/C4/C5/C6/N2/N3) and  $p3$  (C9/C10/  
C11/C12/C13/C14). The dihedral angles between the planes  
are  $0.4$  ( $3$ )° for  $p1/p2$ ,  $134.7$  ( $2$ )° for  $p1/p3$  and  $134.6$  ( $3$ )° for  
 $p2/p3$ .

## Experimental

1-Phenylbutane-1,3-dione (0.32 g, 2 mmol) in ethanol (5 ml) was  
added to a solution of 5-amino-3-methylsulfanyl-1*H*-pyrazole-4-  
carbonitrile (0.31 g, 2 mmol) in ethanol (15 ml) containing a few  
drops of acetic acid. The mixture was refluxed for 5 h then cooled to  
room temperature, and the crude product (0.5 g, 89.3%) was obtained  
without further purification. The isomers were separated by column  
chromatography on silica gel (petroleum ether/ethyl acetate = 3/1  
(*v/v*)) and the ratio of the isomers was 59.2:40.8 for (I):(II).

## Crystal data

 $\text{C}_{15}\text{H}_{12}\text{N}_4\text{S}$   
 $M_r = 280.35$   
Monoclinic,  $P2_1/c$   
 $a = 8.819$  (3) Å  
 $b = 7.822$  (3) Å  
 $c = 20.465$  (8) Å  
 $\beta = 96.507$  (7)°  
 $V = 1402.6$  (9) Å<sup>3</sup>  
 $Z = 4$  $D_x = 1.328$  Mg m<sup>-3</sup>  
Mo  $K\alpha$  radiation  
Cell parameters from 778  
reflections  
 $\theta = 3.2$ – $23.2$ °  
 $\mu = 0.23$  mm<sup>-1</sup>  
 $T = 293$  (2) K  
Block, colourless  
 $0.24 \times 0.22 \times 0.18$  mm

## Data collection

Bruker SMART CCD area-detector diffractometer	2880 independent reflections
$\varphi$ and $\omega$ scans	1655 reflections with $I > 2\sigma(I)$
Absorption correction: multi-scan (SADABS; Sheldrick, 1996)	$R_{\text{int}} = 0.047$
$T_{\text{min}} = 0.863$ , $T_{\text{max}} = 1.000$	$\theta_{\text{max}} = 26.5^\circ$
7758 measured reflections	$h = -11 \rightarrow 9$
	$k = -9 \rightarrow 9$
	$l = -25 \rightarrow 25$

## Refinement

Refinement on $F^2$	$w = 1/[\sigma^2(F_o^2) + (0.0562P)^2 + 0.0043P]$
$R[F^2 > 2\sigma(F^2)] = 0.051$	where $P = (F_o^2 + 2F_c^2)/3$
$wR(F^2) = 0.122$	$(\Delta/\sigma)_{\text{max}} = 0.001$
$S = 1.00$	$\Delta\rho_{\text{max}} = 0.23 \text{ e } \text{\AA}^{-3}$
2880 reflections	$\Delta\rho_{\text{min}} = -0.24 \text{ e } \text{\AA}^{-3}$
183 parameters	
H-atom parameters constrained	

Table 1

Selected geometric parameters ( $\text{\AA}$ ,  $^\circ$ ).

S1—C1	1.735 (3)	N4—C7	1.139 (3)
N1—N2	1.376 (3)	C2—C7	1.409 (4)
N2—C4	1.367 (3)	C4—C5	1.363 (3)
N2—C3	1.381 (3)	C4—C9	1.467 (3)
C1—S1—C8	101.26 (13)	C5—C4—N2	114.7 (2)
N1—N2—C3	112.53 (19)	N4—C7—C2	179.1 (3)
C1—N1—N2—C3	−1.0 (2)	C5—C4—C9—C14	−43.9 (4)

All H atoms were placed in calculated positions, with C—H = 0.93 or 0.96  $\text{\AA}$ , and refined using a riding model, with  $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$ .

Data collection: SMART (Bruker, 1998); cell refinement: SMART; data reduction: SAINT (Bruker, 1999); program(s) used to solve structure: SHELXS97 (Sheldrick, 1997); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: SHELXTL (Bruker, 1999); software used to prepare material for publication: SHELXTL.

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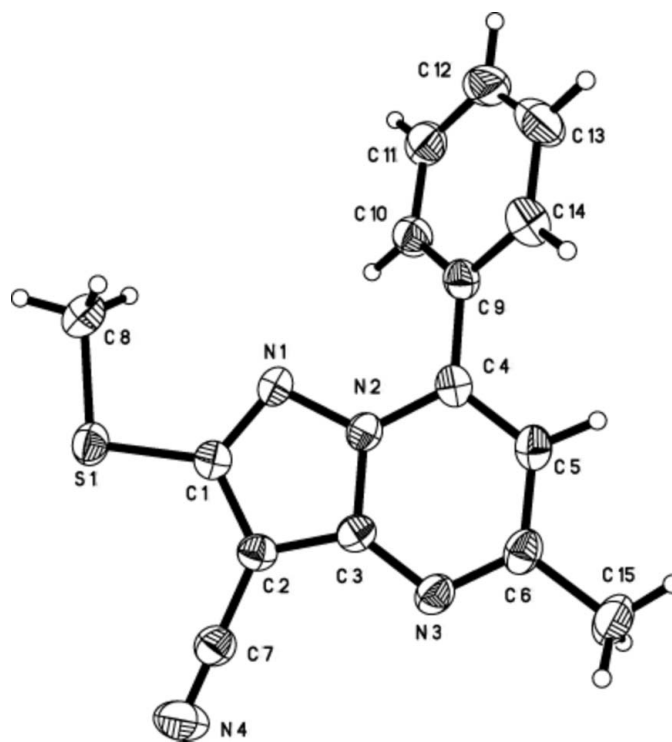


Figure 1

A view of the title compound, showing the atom-numbering scheme. Displacement ellipsoids are drawn at the 50% probability level.

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