

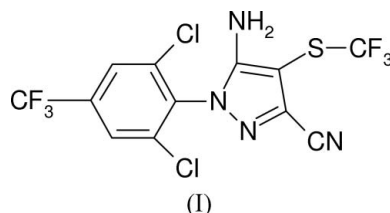
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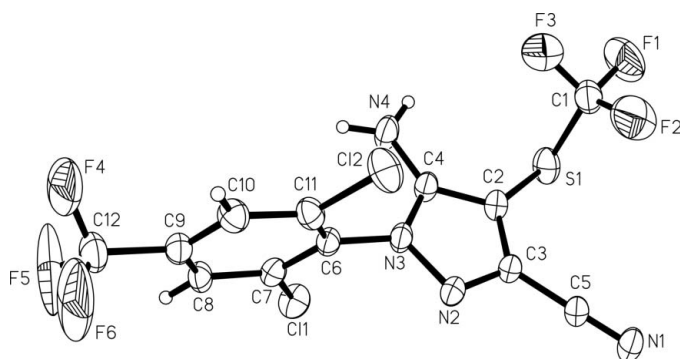
## Key indicators

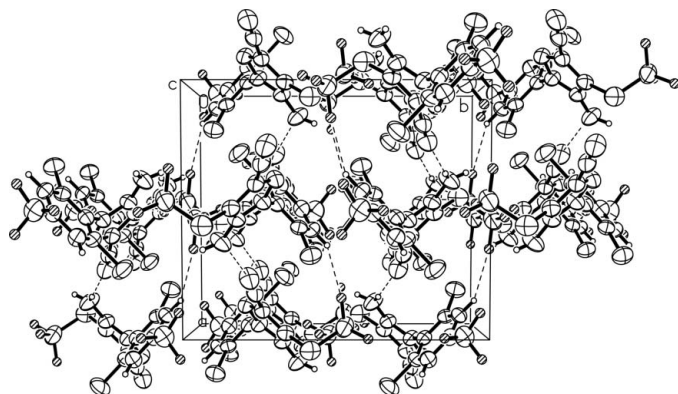
Single-crystal X-ray study  
 $T = 298$  K  
Mean  $\sigma(\text{C}-\text{C}) = 0.003$  Å  
Disorder in main residue  
 $R$  factor = 0.047  
 $wR$  factor = 0.113  
Data-to-parameter ratio = 10.7For details of how these key indicators were  
automatically derived from the article, see  
<http://journals.iucr.org/e>.5-Amino-1-[2,6-dichloro-4-(trifluoromethyl)-  
phenyl]-4-(trifluoromethylsulfanyl)-1*H*-pyrazole-  
3-carbonitrileThe molecule of the title compound,  $\text{C}_{12}\text{H}_4\text{Cl}_2\text{F}_6\text{N}_4\text{S}$ , exhibits  
disorder of both trifluoromethyl groups. Crystal packing is  
realised by  $\text{N}-\text{H} \cdots \text{N}$  hydrogen bonds.Received 3 October 2005  
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## Comment

The title compound, (I) (Fig. 1), is an important starting  
material for the synthesis of fipronil, a very useful insecticide  
(Clavel *et al.*, 2003; Hatton *et al.*, 1993).One of the two trifluoromethyl groups occurs in two equally  
populated orientations, whereas the other oscillates about the  
 $\text{C1}-\text{F1}$  bond, having two equally populated positions for  
atoms F2 and F3. The pyrazole ring and benzene ring make a  
dihedral angle of  $83.15(12)^\circ$ .The regular self-assembly of the title compound (Fig. 2) is  
realised by intermolecular hydrogen bonds: an amino group  
forms a hydrogen bond with the cyano N atom of another  
molecule (Table 2).

## Experimental

According to the method of Wakselman *et al.* (1992), the following  
procedure was employed. Into a 100 ml Teflon-coated autoclave, a  
dimethylformamide solution (48 ml) of 5-amino-3-cyano-1-[2,6-  
dichloro-4-(trifluoromethyl)phenyl]-4-pyrazolyl disulfide (0.8 g) was  
placed, followed by an aqueous solution (24 ml) of sodium hydrogen  
phosphate (1.2 g). Sodium dithionate (0.6 g) was then added with**Figure 1**  
The structure of (I), showing the atom-numbering scheme and with  
displacement ellipsoids at the 50% probability level.



**Figure 2**  
The crystal packing of (I), with N—H...N hydrogen bonds shown as dashed lines.

stirring. The preparation of the disulfide was carried out according to Clavel *et al.* (2003) and Tang *et al.* (2005). Next,  $\text{CF}_3\text{Br}$  was introduced at a pressure of 5–6 bar (autogenous pressure; 1 bar = 100 000 Pa) and, after 5 h of vigorous stirring at room temperature, the title compound was obtained in 51% yield. Single crystals of (I) suitable for X-ray analysis were obtained by slow evaporation of a toluene solution (m.p. 434–436 K). Spectroscopic analysis: IR (KBr,  $\nu$ ,  $\text{cm}^{-1}$ ): 3475, 3331, 3233, 2258, 1631, 1561, 1513, 1463, 1399, 1318, 1153, 1109, 886, 812, 635;  $^1\text{H}$  NMR ( $\text{C}_3\text{D}_6\text{O}$ ,  $\delta$ , p.p.m.): 8.11 (s, 2H), 6.64 (s, 2H);  $^{13}\text{C}$  NMR ( $\text{C}_3\text{D}_6\text{O}$ ,  $\delta$ , p.p.m.): 153.4 (1C), 136.3 (1C), 135.4 (1C), 133.9 (1C), 132.4 (1C), 128.9 (1C), 126.5 (2C), 126.3 (2C), 122.2 (1C), 111.6 (1C);  $^{19}\text{F}$  NMR ( $\text{C}_3\text{D}_6\text{O}$ ,  $\delta$ , p.p.m.):  $-46.25$  (3F),  $-63.63$  (3F).

#### Crystal data

$\text{C}_{12}\text{H}_4\text{Cl}_2\text{F}_6\text{N}_4\text{S}$   
 $M_r = 421.16$   
Monoclinic,  $P2_1/n$   
 $a = 10.7801$  (8) Å  
 $b = 12.7006$  (9) Å  
 $c = 12.1039$  (9) Å  
 $\beta = 96.973$  (1)°  
 $V = 1644.9$  (2) Å<sup>3</sup>  
 $Z = 4$

$D_x = 1.701$  Mg m<sup>-3</sup>  
Mo  $K\alpha$  radiation  
Cell parameters from 3781 reflections  
 $\theta = 2.3$ – $25.0^\circ$   
 $\mu = 0.59$  mm<sup>-1</sup>  
 $T = 298$  (2) K  
Prism, colourless  
 $0.42 \times 0.29 \times 0.21$  mm

#### Data collection

Bruker APEX area-detector diffractometer  
 $\varphi$  and  $\omega$  scans  
Absorption correction: multi-scan (SADABS; Bruker, 2002)  
 $T_{\min} = 0.791$ ,  $T_{\max} = 0.887$   
8518 measured reflections

2953 independent reflections  
2636 reflections with  $I > 2\sigma(I)$   
 $R_{\text{int}} = 0.018$   
 $\theta_{\text{max}} = 25.2^\circ$   
 $h = -12 \rightarrow 12$   
 $k = -12 \rightarrow 15$   
 $l = -14 \rightarrow 12$

#### Refinement

Refinement on  $F^2$   
 $R[F^2 > 2\sigma(F^2)] = 0.047$   
 $wR(F^2) = 0.114$   
 $S = 1.10$   
2953 reflections  
277 parameters  
H atoms treated by a mixture of independent and constrained refinement

$w = 1/[\sigma^2(F_o^2) + (0.059P)^2 + 0.9497P]$   
where  $P = (F_o^2 + 2F_c^2)/3$   
 $(\Delta/\sigma)_{\text{max}} = 0.001$   
 $\Delta\rho_{\text{max}} = 0.21$  e Å<sup>-3</sup>  
 $\Delta\rho_{\text{min}} = -0.26$  e Å<sup>-3</sup>

**Table 1**  
Selected geometric parameters (Å, °).

C11—C7	1.726 (2)	N3—C4	1.349 (3)
C12—C11	1.727 (3)	N3—C6	1.420 (3)
S1—C2	1.735 (3)	N4—C4	1.340 (3)
S1—C1	1.783 (4)	C3—C5	1.438 (3)
F3—C1	1.266 (11)	C6—C11	1.383 (4)
F4—C12	1.321 (8)	C6—C7	1.384 (3)
N1—C5	1.135 (3)	C8—C9	1.377 (4)
N2—C3	1.323 (3)	C8—H8	0.9300
N2—N3	1.374 (3)	C9—C12	1.503 (4)
C2—S1—C1	100.16 (15)	N1—C5—C3	179.0 (3)
C3—N2—N3	102.7 (2)	C11—C6—C7	119.0 (2)
C4—N3—N2	113.58 (19)	C11—C6—N3	120.4 (2)
C4—N3—C6	126.99 (19)	C7—C6—N3	120.6 (2)
N2—N3—C6	119.44 (19)	C8—C7—C6	121.1 (2)
H4A—N4—H4B	123 (3)	C8—C7—C11	119.27 (19)
F3—C1—F2	111.0 (13)	C9—C8—H8	120.7
F3—C1—F1	110.7 (7)	C10—C9—C8	121.6 (2)
F2—C1—F1	100.7 (13)	C10—C9—C12	119.1 (3)
C4—C2—C3	104.4 (2)	C10—C11—C6	120.7 (2)
C4—C2—S1	125.86 (19)	F5—C12—F4	71.4 (7)
C3—C2—S1	129.47 (18)	F6'—C12—F6	48.2 (6)
N4—C4—N3	123.2 (2)	F5—C12—C9	112.3 (6)
N4—C4—C2	130.9 (2)		

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

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
$\text{N4—H4A}\cdots\text{N1}^i$	0.85 (2)	2.27 (2)	3.099 (3)	164 (3)

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

All H atoms were placed in geometrically idealized positions and constrained to ride on their parent atoms, with N—H = 0.83–0.85 Å and C—H = 0.93 Å, and with  $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C}, \text{N})$ . The two  $\text{CF}_3$  groups were treated as disordered over two positions using free variables.

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

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