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Key indicators

Single-crystal X-ray study
T = 293 K
Mean $\sigma(\text{C}-\text{C}) = 0.002 \text{ \AA}$
Disorder in main residue
R factor = 0.030
wR factor = 0.085
Data-to-parameter ratio = 14.9

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

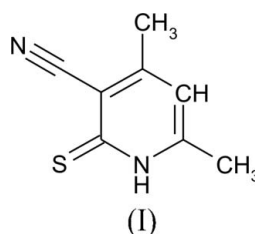
4,6-Dimethyl-2-thioxo-1,2-dihydropyridine-3-carbonitrile

In the crystal structure of the title compound, $\text{C}_8\text{H}_8\text{N}_2\text{S}$, the molecules form centrosymmetric dimers *via* $\text{N}-\text{H}\cdots\text{S}$ hydrogen bonds.

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Comment

3-Cyanopyridine-2(1*H*)-thiones have attracted great attention from many researchers in the past 10–15 years, due to their unusual properties and possibilities for use as synthons for preparing novel biologically active compounds (Litvinov *et al.*, 1989, 1992). Over this period of time, the chemistry of 3-cyanopyridine-2(1*H*)-thione and its derivatives has been extensively studied, including the modification of the nitrile group by hydrolysis (Zhiryakov & Abramenko, 1965; Jahine *et al.*, 1979), reactions with organometallic compounds, and intra- and intermolecular heterocyclization (Litvinov *et al.*, 1989, 1992). Here, we report the crystal structure of the title compound, (I).

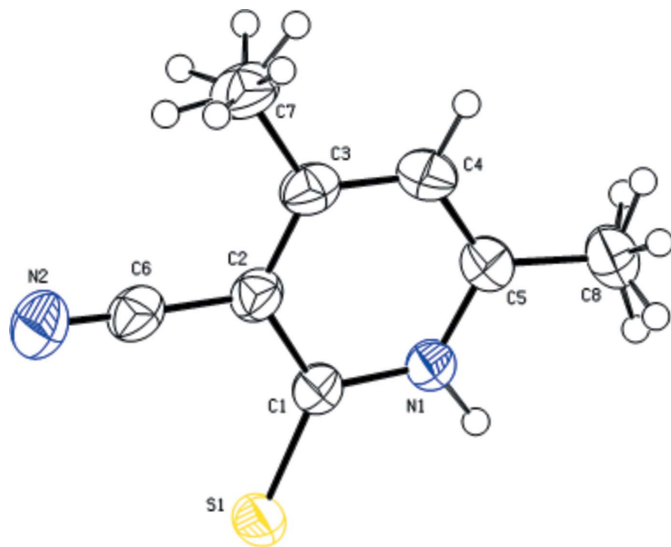


The molecular structure of (I) is shown in Fig. 1, with the atom-numbering scheme. The pyridine ring in (I) is planar. The six-membered heterocycle has a well defined diene-like structure. The C2–C3 and C4–C5 bonds are shorter than the C1–C2 and C3–C4 bonds. The rigid cyano group has the standard linear structure and the C6–N2 bond distance is 1.140 (2) Å. The C1–S1 bond distance is 1.6808 (14) Å. The C–C bond lengths to the methyl groups are C3–C7 = 1.499 (2) and C5–C8 = 1.487 (2) Å. The other bond lengths and angles in (I) are within expected ranges and are similar to those reported in other studies (Rybakov *et al.*, 2004; Zavodnik *et al.*, 2005; Begum & Vasundhara, 2006; Albov *et al.*, 2004).

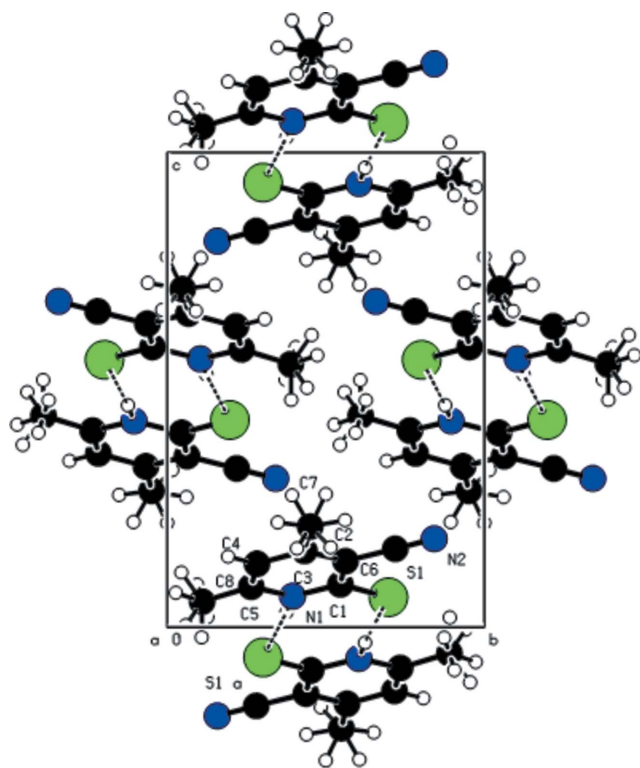
An $\text{N}-\text{H}\cdots\text{S}$ intermolecular hydrogen bond links the molecules in the crystal structure into centrosymmetric dimers (Fig. 2, Table 2).

Experimental

A mixture of 2-cyanothioacetamide (0.10 g, 0.01 mol), acetylacetone (1.03 ml, 1 g, 0.01 mol) and methanol (10 ml) was warmed; to the solution KOH (0.72 g) in methanol (50 ml) was added with stirring


Figure 1

A view of the molecular structure of (I), with the atom-numbering scheme. Displacement ellipsoids are drawn at the 50% probability level. Both disorder components are shown.


Figure 2

The packing of (I). Dashed lines indicate hydrogen bonds. Both disorder components are shown.

for 2 h. During the addition, a coloured precipitate formed and sufficient methanol was added to prevent caking. The mixture was heated at reflux temperature and stirred for 4 h, then cooled to room temperature and acidified with dilute HCl. The product was removed from the cooled reaction mixture by filtration, washed with water, dried, and recrystallized from methanol.

Crystal data

$C_8H_8N_2S$
 $M_r = 164.22$
 Monoclinic, $P2_1/c$
 $a = 7.6526$ (6) Å
 $b = 8.4577$ (4) Å
 $c = 13.7386$ (11) Å
 $\beta = 113.185$ (6)°

$V = 817.39$ (10) Å³
 $Z = 4$
 Mo $K\alpha$ radiation
 $\mu = 0.33$ mm⁻¹
 $T = 293$ (2) K
 $0.44 \times 0.34 \times 0.19$ mm

Data collection

Stoe IPDS II diffractometer
 Absorption correction: integration
 (*X-RED32*; Stoe & Cie, 2002)
 $T_{\min} = 0.856$, $T_{\max} = 0.939$

11418 measured reflections
 1606 independent reflections
 1431 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.031$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.030$
 $wR(F^2) = 0.085$
 $S = 1.06$
 1606 reflections
 108 parameters

H atoms treated by a mixture of independent and constrained refinement
 $\Delta\rho_{\max} = 0.18$ e Å⁻³
 $\Delta\rho_{\min} = -0.21$ e Å⁻³

Table 1

Selected geometric parameters (Å, °).

C1—N1	1.3593 (17)	C5—N1	1.3509 (18)
C1—S1	1.6808 (14)		
N1—C1—C2	114.79 (12)	N1—C5—C8	117.13 (13)
N1—C1—S1	120.78 (10)	N2—C6—C2	177.89 (16)
N1—C5—C4	118.69 (14)	C5—N1—C1	125.59 (12)

Table 2

Hydrogen-bond geometry (Å, °).

<i>D</i> —H... <i>A</i>	<i>D</i> —H	H... <i>A</i>	<i>D</i> ... <i>A</i>	<i>D</i> —H... <i>A</i>
N1—H1...S1 ⁱ	0.925 (19)	2.36 (2)	3.2808 (13)	172.9 (17)

Symmetry code: (i) $-x, -y + 1, -z$.

The H atoms bonded to C4 and N1 were located in a difference Fourier synthesis and refined freely [$C-H = 0.95$ (2) Å]. The methyl H atoms were positioned geometrically, with $C-H = 0.96$ Å and $U_{\text{iso}}(H) = 1.5U_{\text{eq}}(C)$. The methyl group is disordered over two positions with equal occupancy.

Data collection: *X-AREA* (Stoe & Cie, 2002); cell refinement: *X-AREA*; data reduction: *X-RED32* (Stoe & Cie, 2002); program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *ORTEP-3* (Farrugia, 1997); software used to prepare material for publication: *WinGX* (Farrugia, 1999).

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