

## 2-Amino-4-methylthiophene-3-carbonitrile

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## Key indicators

Single-crystal X-ray study

$T = 293\text{ K}$

Mean  $\sigma(\text{C}-\text{C}) = 0.005\text{ \AA}$

$R$  factor = 0.052

$wR$  factor = 0.154

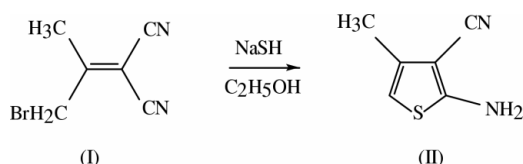
Data-to-parameter ratio = 13.8

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

The title compound,  $\text{C}_6\text{H}_6\text{N}_2\text{S}$ , consists of a thiophene ring carrying three substituent groups. Two  $\text{N}-\text{H}\cdots\text{N}$  intermolecular hydrogen bonds link neighboring molecules into a three-dimensional network.

## Comment

Highly substituted thiophenes form an important part of numerous natural products (Koike *et al.*, 1997) and pharmaceuticals (World Drug Index, 2000). They are often used as novel conducting polymers and isostatic replacements for phenyl groups in medicinal chemistry (Jarvest *et al.*, 1999). The electronic and optical properties of polythiophene and its derivatives have been the subject of many investigations (Roncali, 1997; Ekinci & Demir, 2002).



The title compound, (II) (Fig. 1), consists of a thiophene ring carrying three substituents, *viz.* an amino group, a methyl group and a cyano group. The  $\text{S}-\text{C}$  bond lengths, 1.728 (4) and 1.725 (4)  $\text{\AA}$ , are in good agreement with those in the literature, *e.g.* 1.734 (2) and 1.721 (3)  $\text{\AA}$  (Han & Choi, 2000), 1.727 (1) and 1.729 (2)  $\text{\AA}$  (Elerman & Elmalı, 1998), and 1.723 (2) and 1.735 (3)  $\text{\AA}$  (Wouters *et al.*, 1997). The  $\text{C1}\equiv\text{N1}$  bond distance is 1.153 (4)  $\text{\AA}$ , typical of such a triple bond. This value agrees well with similar bonds reported in the literature, *e.g.* 1.132 (2)  $\text{\AA}$  (Elerman & Elmalı, 1998), 1.130 (5) and 1.142 (5)  $\text{\AA}$  (Çoruh *et al.*, 2002), and 1.148 (2)  $\text{\AA}$  (Boitsov *et al.*, 2002). The molecule of (II) is planar, the maximum deviation from the least-squares plane being 0.0074 (33)  $\text{\AA}$  for atom C4.

A packing diagram of (II) is shown Fig. 2. The molecules are stacked such that there are no ring–ring or  $X-\text{H}\cdots\pi$  interactions between molecules in the stacks; the distance between planes of molecules in the stacks is 5.671 (7)  $\text{\AA}$ . The crystal structure of (II) is stabilized by two  $\text{N}-\text{H}\cdots\text{N}$  intermolecular hydrogen bonds (Table 3 and Fig. 2).

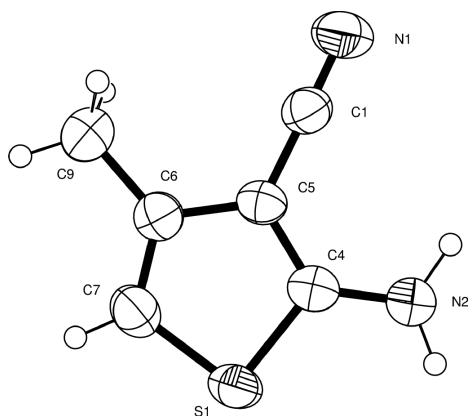
## Experimental

2-(2-Bromo-1-methylethylidene)malonitrile (0.94 g, 5 mmol) was dissolved in a solution of dioxane (5 ml) and absolute ethanol (20 ml). The stirred solution was cooled to 273 K in an ice-salt bath, and a suspension of NaSH (0.3 g) in 10 ml of absolute ethanol was then added dropwise over a period of 30 min. The resulting reaction

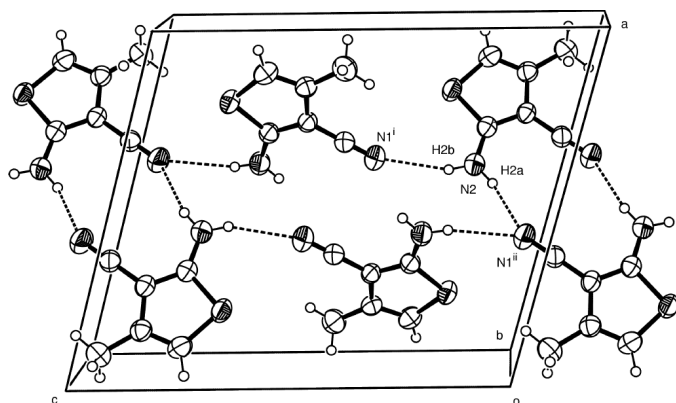
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**Figure 1**  
View of the title compound, with the atom numbering scheme. Displacement ellipsoids for non-H atoms are drawn at the 50% probability level.



**Figure 2**  
The hydrogen-bond network in (II), viewed approximately along the [010] direction. Hydrogen bonds are indicated by dashed lines.

mixture was stirred for a further 1 h at room temperature. After removal of the solvent, the residue was filtered on a short  $\text{Al}_2\text{O}_3$  column, eluting with hexane–ethyl acetate (7:3). The solvent was removed and the residue crystallized from chloroform to yield (556 mg, 81%) 2-amino-4-methylthiophene-3-carbonitrile (light pink crystals, mp 391–392 K).  $^1\text{H}$  NMR (200 MHz,  $\text{CDCl}_3$ ):  $\delta$  5.59 (s, 1H, H<sub>5</sub>), 4.81 (bs, 2H, NH<sub>2</sub>), 2.11 (s, 3H, CH<sub>3</sub>);  $^{13}\text{C}$  NMR (50 MHz,  $\text{CDCl}_3$ )  $\delta$  162.44, 135.91, 115.52, 105.32, 91.03, 15.53. IR ( $\text{CHCl}_3$ ) 3417, 3212, 3097, 2977, 2202, 1627, 1558, 1446, 1300, 1191, 1068, 836.

#### Crystal data

$\text{C}_6\text{H}_6\text{N}_2\text{S}$	$D_x = 1.376 \text{ Mg m}^{-3}$
$M_r = 138.20$	Mo $K\alpha$ radiation
Monoclinic, $P2_1/c$	Cell parameters from 1153 reflections
$a = 11.900$ (5) Å	$\theta = 1.8\text{--}25.2^\circ$
$b = 4.130$ (5) Å	$\mu = 0.39 \text{ mm}^{-1}$
$c = 14.085$ (5) Å	$T = 293 \text{ K}$
$\beta = 105.562$ (5)°	Block, pink
$V = 666.9$ (9) Å <sup>3</sup>	$0.20 \times 0.15 \times 0.10 \text{ mm}$
$Z = 4$	

#### Data collection

Nonius KappaCCD diffractometer	$R_{\text{int}} = 0.060$
$\varphi$ and $\omega$ scans	$\theta_{\text{max}} = 25.2^\circ$
Absorption correction: none	$h = -13 \rightarrow 13$
1849 measured reflections	$k = -4 \rightarrow 4$
1135 independent reflections	$l = -16 \rightarrow 16$
789 reflections with $I > 2\sigma(I)$	

#### Refinement

Refinement on $F^2$	H-atom parameters constrained
$R[F^2 > 2\sigma(F^2)] = 0.052$	$w = 1/[\sigma^2(F_o^2) + (0.087P)^2]$
$wR(F^2) = 0.154$	where $P = (F_o^2 + 2F_c^2)/3$
$S = 1.00$	$(\Delta/\sigma)_{\text{max}} < 0.001$
1135 reflections	$\Delta\rho_{\text{max}} = 0.33 \text{ e \AA}^{-3}$
82 parameters	$\Delta\rho_{\text{min}} = -0.31 \text{ e \AA}^{-3}$

**Table 1**

Selected geometric parameters (Å, °).

S1–C4	1.728 (4)	N1–C1	1.153 (4)
S1–C7	1.725 (4)	N2–C4	1.354 (5)
C4–S1–C7	91.95 (15)	S1–C4–C5	110.3 (3)
N1–C1–C5	179.2 (4)	N2–C4–C5	128.8 (3)
S1–C4–N2	120.9 (2)	S1–C7–C6	112.9 (3)

**Table 2**

Contact distances (Å, °).

S1...N1 <sup>i</sup>	3.401 (5)	N1...N2 <sup>ii</sup>	3.117 (6)
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**Table 3**

Hydrogen-bonding geometry (Å, °).

$D\text{--}H\cdots A$	$D\text{--}H$	$H\cdots A$	$D\cdots A$	$D\text{--}H\cdots A$
N2–H2B...N1 <sup>i</sup>	0.86	2.48	3.276 (5)	155
N2–H2A...N1 <sup>ii</sup>	0.86	2.31	3.117 (6)	157

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

All H atoms were positioned geometrically and refined in riding mode, with methyl C–H = 0.96 Å, other C–H = 0.93 Å, and N–H = 0.86 Å. For methyl H,  $U_{\text{iso}}$  values were set equal to  $1.5U_{\text{eq}}$  of the carrier C atom. For other N–H and other C–H,  $U_{\text{iso}}(\text{H})$  values were set equal to  $1.2U_{\text{eq}}$  of the carrier atom.

Data collection: COLLECT (Nonius BV, 1997–2000); cell refinement: HKL SCALEPACK (Otwinowski & Minor, 1997); data reduction: HKL DENZO (Otwinowski & Minor, 1997) and SCALEPACK; program(s) used to solve structure: SHELXS97 (Sheldrick, 1997); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: ORTEP-3 for Windows (Farrugia, 1997) and PLATON (Spek, 1997); software used to prepare material for publication: SHELXL97 and WinGX (Farrugia, 1999).

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