

7-Amino-1*H*-3,1-benzothiazine-2,4-dithioneYun Yu,^a Hui-Ping Zhong,^b
Kai-Bin Yang,^b Rong-Bin
Huang^{b*} and Lan-Sun Zheng^c^aDepartment of Chemistry, Long Yan College, Long Yan, Fujian 361005, People's Republic of China, ^bDepartment of Chemistry, Xiamen University, Xiamen 361005, People's Republic of China, and ^cState Key Laboratory for Physical Chemistry of Solid Surfaces, Xiamen University, Xiamen 361005, People's Republic of China

Correspondence e-mail: rbhuang@xmu.edu.cn

Key indicators

Single-crystal X-ray study

 $T = 273\text{ K}$ Mean $\sigma(\text{C}-\text{C}) = 0.004\text{ \AA}$ R factor = 0.059 wR factor = 0.160

Data-to-parameter ratio = 17.3

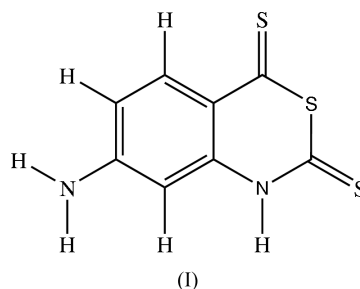
For details of how these key indicators were automatically derived from the article, see <http://journals.iucr.org/e>.X-ray crystal structure determination shows that the title compound, $\text{C}_8\text{H}_6\text{N}_2\text{S}_3$, has a highly planar molecule, packing *via* amino–thione hydrogen bonds.

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Comment

The chemistry of thiocarbonyl (thione) analogues of such carbonyl functions as ketones, acids, esters and amides has been the subject of extensive study (Duus, 1979). Lakshmi-kantham *et al.* (1984) reported the crystal structure of the first thione analogue of a cyclic anhydride. We report here the synthesis and characterization of a new thione analogue of a cyclic anhydride containing nitrogen, 7-amino-1*H*-benzo[*d*]-[1,3]thiazine-2,4-dithione, (I), obtained by the hydrothermal reaction of 1,3-phenylenediamine and carbon disulfide.

The present crystal structure analysis shows that the molecule of (I) is highly planar, with a small (0.0054 Å) mean

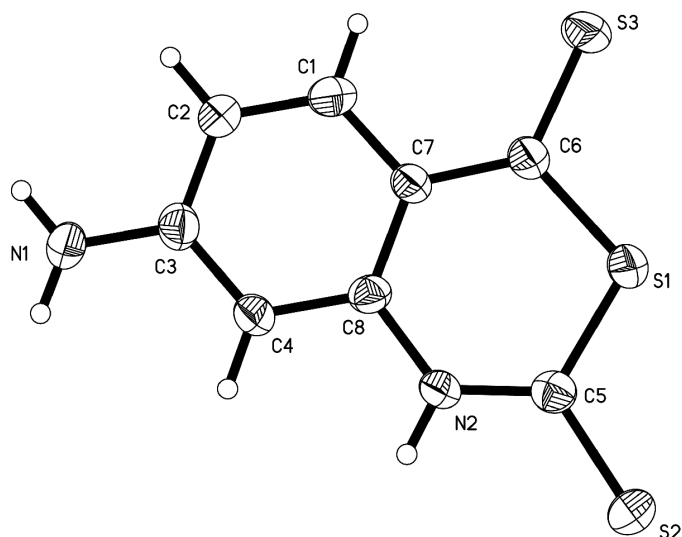


Figure 1

A view of the molecule of (I), with the atom-numbering scheme. Displacement ellipsoids are drawn at 50% probability level and H atoms are shown as small spheres of arbitrary radii.

deviation from the plane including all non-H atoms. The molecule features long S1—C5 [1.727 (3) Å] and S1—C6 [1.735 (3) Å] bonds, and short S2—C5 [1.663 (3) Å] and S3—C6 [1.660 (3) Å] bonds.

The molecules of (I) pack in the solid state *via* amino-thione hydrogen bonds, with N2···S3ⁱ 3.407 (3) Å, N1···S2ⁱⁱ 3.530 (3) Å and N1···S2ⁱⁱⁱ 3.610 (3) Å [symmetry codes: (i) $x, \frac{1}{2} - y, z + \frac{1}{2}$; (ii) $1 + x, \frac{1}{2} - y, z + \frac{1}{2}$; (iii) $1 - x, y - \frac{1}{2}, \frac{3}{2} - z$; Table 2).

Experimental

An ethanol solution (20 ml) containing *m*-phenylenediamine dihydrochloride (5 mmol) and carbon disulfide (10 mmol) was placed in a 25 ml autoclave with a Teflon liner. The pH of the solution was adjusted to 7–8 with sodium hydroxide solution. The autoclave was heated to 373 K, maintained at that temperature for 96 h and cooled to room temperature at a rate of 0.5 K min⁻¹. Red block-shaped crystals of (I) were obtained for X-ray diffraction. Analysis, calculated for C₈H₆N₂S₃: C 42.45, H 2.67, N 12.38, S 42.50%; found: C 43.16, H 2.55, N 12.67, S 41.97%. Spectroscopic analysis: ¹H NMR (CD₃COCD₃, δ, p.p.m.): 6.65 (ArH), 6.68 (NH₂), 6.73 (ArH), 8.25 (ArH), 12.04 (NH).

Crystal data

C ₈ H ₆ N ₂ S ₃	$D_x = 1.642 \text{ Mg m}^{-3}$
$M_r = 226.33$	Mo $K\alpha$ radiation
Monoclinic, $P2_1/c$	Cell parameters from 4962 reflections
$a = 3.9117 (9) \text{ \AA}$	$\theta = 2.0\text{--}27.0^\circ$
$b = 17.232 (4) \text{ \AA}$	$\mu = 0.76 \text{ mm}^{-1}$
$c = 13.686 (3) \text{ \AA}$	$T = 273 (2) \text{ K}$
$\beta = 97.116 (4)^\circ$	Block, red
$V = 915.4 (3) \text{ \AA}^3$	$0.38 \times 0.22 \times 0.16 \text{ mm}$
$Z = 4$	

Data collection

Bruker SMART APEX CCD area-detector diffractometer	2042 independent reflections
φ and ω scans	1779 reflections with $I > 2\sigma(I)$
Absorption correction: multi-scan (SADABS; Sheldrick, 1996)	$R_{\text{int}} = 0.028$
$T_{\text{min}} = 0.762, T_{\text{max}} = 0.889$	$\theta_{\text{max}} = 27.5^\circ$
5279 measured reflections	$h = -4 \rightarrow 5$
	$k = -17 \rightarrow 22$
	$l = -17 \rightarrow 15$

Refinement

Refinement on F^2	$w = 1/[\sigma^2(F_o^2) + (0.0963P)^2 + 0.2434P]$
$R[F^2 > 2\sigma(F^2)] = 0.059$	where $P = (F_o^2 + 2F_c^2)/3$
$wR(F^2) = 0.160$	$(\Delta/\sigma)_{\text{max}} < 0.0001$
$S = 1.14$	$\Delta\rho_{\text{max}} = 0.75 \text{ e \AA}^{-3}$
2042 reflections	$\Delta\rho_{\text{min}} = -0.31 \text{ e \AA}^{-3}$
118 parameters	
H-atom parameters constrained	

Table 1

Selected geometric parameters (Å, °).

S1—C5	1.727 (3)	C7—C8	1.409 (4)
S1—C6	1.735 (3)	C7—C6	1.431 (4)
S2—C5	1.663 (3)	C4—C3	1.388 (4)
S3—C6	1.660 (3)	C4—C8	1.393 (4)
N2—C5	1.338 (4)	N1—C3	1.359 (4)
N2—C8	1.396 (4)	C1—C2	1.366 (4)
C7—C1	1.405 (4)	C3—C2	1.409 (4)
C5—S1—C6	107.22 (14)	C7—C6—S1	119.8 (2)
C5—N2—C8	127.8 (2)	S3—C6—S1	113.21 (17)
C1—C7—C8	117.6 (3)	C2—C1—C7	122.1 (3)
C1—C7—C6	119.8 (3)	N2—C8—C4	116.4 (2)
C8—C7—C6	122.7 (3)	N2—C8—C7	123.0 (2)
C3—C4—C8	120.6 (3)	C4—C8—C7	120.6 (3)
N2—C5—S2	123.8 (2)	N1—C3—C4	121.0 (3)
N2—C5—S1	119.5 (2)	N1—C3—C2	119.8 (3)
S2—C5—S1	116.64 (17)	C4—C3—C2	119.1 (3)
C7—C6—S3	127.0 (2)	C1—C2—C3	120.0 (3)

Table 2

Hydrogen-bond geometry (Å, °).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
N2—H2B···S3 ⁱ	0.86	2.58	3.407 (3)	163
N1—H1B···S2 ⁱⁱ	0.86	2.72	3.530 (3)	157
N1—H1C···S2 ⁱⁱⁱ	0.86	2.79	3.610 (3)	160
Symmetry codes:	(i) $x, -y + \frac{1}{2}, z + \frac{1}{2}$	(ii) $x + 1, -y + \frac{1}{2}, z + \frac{1}{2}$	(iii) $-x + 1, y - \frac{1}{2}, -z + \frac{3}{2}$	

All H atoms were positioned geometrically, with C—H distances of 0.93 Å and N—H distances of 0.86 Å, and were refined using a riding-atom model, with $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C}, \text{N})$.

Data collection: SMART (Bruker, 2001); cell refinement: SAINT (Bruker, 2001); data reduction: SAINT; program(s) used to solve structure: SHELXS97 (Sheldrick, 1997); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: ORTEPII (Johnson, 1976); software used to prepare material for publication: SHELXL97.

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