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(Z)-2-Sulfanylidene-5-(thiophen-2-yl-methylidene)imidazolidin-4-one

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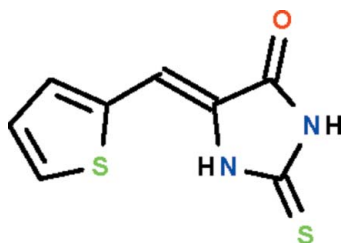
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Key indicators: single-crystal X-ray study; $T = 100$ K; mean $\sigma(\text{C}-\text{C}) = 0.003$ Å; disorder in main residue; R factor = 0.043; wR factor = 0.122; data-to-parameter ratio = 12.5.

The molecule of the title compound, $\text{C}_8\text{H}_6\text{N}_2\text{OS}_2$, has a V shape with two five-membered rings attached to a methylene C atom. All non-H atoms are approximately coplanar (r.m.s. deviation = 0.096 Å). In the crystal, molecules are linked by $\text{N}-\text{H}\cdots\text{O}$ hydrogen bonds into layers. The thiophene ring is disordered over two positions; the major orientation has an occupancy of 0.683 (3). Is there an intramolecular $\text{N}-\text{H}\cdots\text{S}$ bond?

Related literature

For two 5-aryl-2-thioxoimidazolin-4-ones, see: Chowdhry *et al.* (2000); Książek *et al.* (2009).



Experimental

Crystal data

 $\text{C}_8\text{H}_6\text{N}_2\text{OS}_2$
 $M_r = 210.27$

Triclinic, $P\bar{1}$
 $a = 6.1022$ (6) Å
 $b = 7.0806$ (8) Å
 $c = 11.0425$ (13) Å
 $\alpha = 72.582$ (11)°
 $\beta = 76.116$ (10)°
 $\gamma = 75.640$ (9)°

$V = 433.87$ (8) Å³
 $Z = 2$
 Cu $K\alpha$ radiation
 $\mu = 5.22$ mm⁻¹
 $T = 100$ K
 $0.25 \times 0.20 \times 0.02$ mm

Data collection

Agilent SuperNova Dual diffractometer with an Atlas detector
 Absorption correction: multi-scan (*CrysAlis PRO*; Agilent, 2010)
 $T_{\min} = 0.356$, $T_{\max} = 0.903$

2599 measured reflections
 1677 independent reflections
 1519 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.027$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.043$
 $wR(F^2) = 0.122$
 $S = 1.04$
 1677 reflections
 134 parameters

6 restraints
 H-atom parameters constrained
 $\Delta\rho_{\max} = 0.33$ e Å⁻³
 $\Delta\rho_{\min} = -0.44$ e Å⁻³

Table 1

Hydrogen-bond geometry (Å, °).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
$\text{N2}-\text{H2}\cdots\text{O1}^i$	0.88	2.20	2.873 (2)	133

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

Data collection: *CrysAlis PRO* (Agilent, 2010); cell refinement: *CrysAlis PRO*; data reduction: *CrysAlis PRO*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *X-SEED* (Barbour, 2001); software used to prepare material for publication: *publCIF* (Westrip, 2010).

We thank King Abdulaziz University and the University of Malaya for supporting this study.

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: BT5612).

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supporting information

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(Z)-2-Sulfanylidene-5-(thiophen-2-ylmethylidene)imidazolidin-4-one

Abdullah M. Asiri, Hassan M. Faidallah, Abdulrahman O. Al-Youbi, Tarik R. Sobahi and Seik Weng Ng

S1. Comment

The crystal structures of only a small number of 5-aryl-2-thioxoimidazolidin-4-ones have been reported, with that of the phenyl homolog been described only recently. The bond dimensions of the parent compound are used to explain the nature of the products of its cycloaddition reaction (Książek *et al.*, 2009). The 2-pyridyl derivative is also a flat molecule (Chowdhry *et al.*, 2000) The thienyl analog (Scheme I) is similarly planar (r.m.s. deviation 0.096 Å). The molecule has a somewhat butterfly shape with the two five-membered rings attached to the methylene carbon (Fig. 1). The –NH– unit at the 4-position (of one ring) points towards the S atom in the 2-position (of the other ring); however, the interaction is too weak to lock the molecule so that the thienyl ring is able to adopt two orientations. Two molecules are linked by an N–H···O hydrogen bond across a center-of-inversion to form a dimer.

S2. Experimental

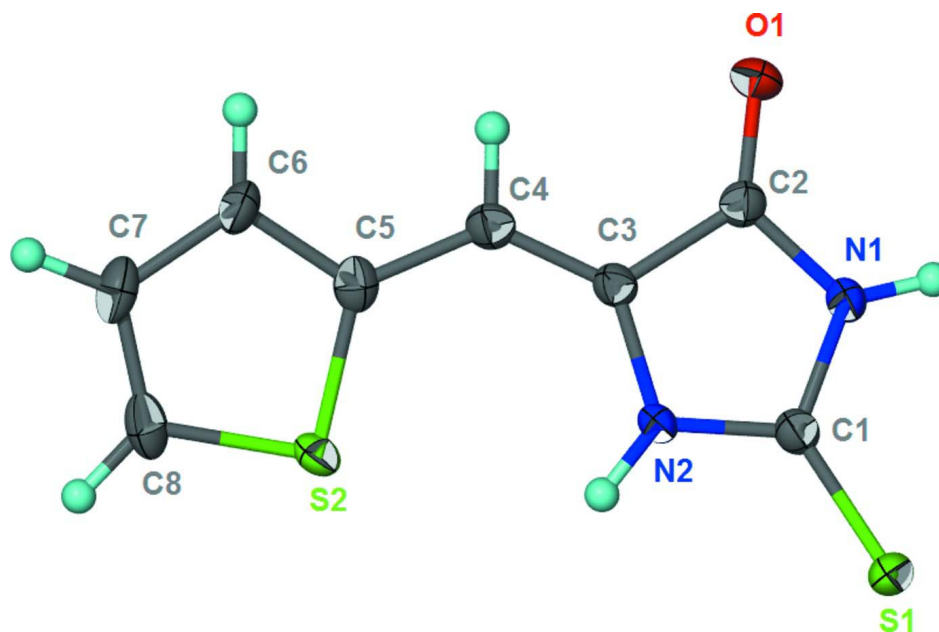
Thiophene-2-carboxaldehyde (1.10 g, 10 mmol) in ethanol (20 ml) was added to a solution of the 2-thiohydantoin (1.16 g, 10 mmol) in 20% ethanolic potassium hydroxide (20 ml). The mixture was stirred for 6 h. The mixture was then poured into water (200 ml). The precipitate that separated when this was acidified with 10% hydrochloric acid was collected and recrystallized from ethanol.

S3. Refinement

H-atoms were placed in calculated positions [C—H 0.95 0.98 and N—H 0.88 Å; $U_{\text{iso}}(\text{H}) 1.2U_{\text{eq}}(\text{C},\text{N})$] and were included in the refinement in the riding model approximation.

The thienyl ring is disordered over two positions; pairs of bond distances were restrained to within 0.01 Å of each other, and the displacement parameters of the overlaying atoms were set to be equal. The major disorder component refined to 68.3 (3)%.

The intensity measurements are complete to 95%; however, they are 100% complete at a 2θ limit of 135 °.

**Figure 1**

Anisotropic displacement ellipsoid plot (Barbour, 2001) of $C_8H_6N_2OS_2$ at the 70% probability level; hydrogen atoms are drawn as spheres of arbitrary radius. The disorder in the thienyl ring is not shown.

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Crystal data

$C_8H_6N_2OS_2$

$M_r = 210.27$

Triclinic, $P\bar{1}$

Hall symbol: -P 1

$a = 6.1022$ (6) Å

$b = 7.0806$ (8) Å

$c = 11.0425$ (13) Å

$\alpha = 72.582$ (11)°

$\beta = 76.116$ (10)°

$\gamma = 75.640$ (9)°

$V = 433.87$ (8) Å³

$Z = 2$

$F(000) = 216$

$D_x = 1.610$ Mg m⁻³

Cu $K\alpha$ radiation, $\lambda = 1.54184$ Å

Cell parameters from 1386 reflections

$\theta = 4.3\text{--}73.9^\circ$

$\mu = 5.22$ mm⁻¹

$T = 100$ K

Plate, yellow

$0.25 \times 0.20 \times 0.02$ mm

Data collection

Agilent SuperNova Dual

diffractometer with an Atlas detector

Radiation source: SuperNova (Cu) X-ray

Source

Mirror monochromator

Detector resolution: 10.4041 pixels mm⁻¹

ω scans

Absorption correction: multi-scan

(*CrysAlis PRO*; Agilent, 2010)

$T_{\min} = 0.356$, $T_{\max} = 0.903$

2599 measured reflections

1677 independent reflections

1519 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.027$

$\theta_{\max} = 74.1^\circ$, $\theta_{\min} = 4.3^\circ$

$h = -4 \rightarrow 7$

$k = -8 \rightarrow 8$

$l = -11 \rightarrow 13$

Refinement

Refinement on F^2

Least-squares matrix: full

$R[F^2 > 2\sigma(F^2)] = 0.043$

$wR(F^2) = 0.122$

$S = 1.04$

1677 reflections

134 parameters

6 restraints

Primary atom site location: structure-invariant
direct methods

Secondary atom site location: difference Fourier
map

Hydrogen site location: inferred from
neighbouring sites

H-atom parameters constrained

$w = 1/[\sigma^2(F_o^2) + (0.0832P)^2 + 0.1141P]$

where $P = (F_o^2 + 2F_c^2)/3$

$(\Delta/\sigma)_{\max} = 0.001$

$\Delta\rho_{\max} = 0.33 \text{ e } \text{\AA}^{-3}$

$\Delta\rho_{\min} = -0.44 \text{ e } \text{\AA}^{-3}$

Fractional atomic coordinates and isotropic or equivalent isotropic displacement parameters (\AA^2)

	x	y	z	$U_{\text{iso}}^*/U_{\text{eq}}$	Occ. (<1)
S1	0.82425 (8)	0.32671 (8)	0.07136 (5)	0.0212 (2)	
S2	0.7731 (2)	0.1352 (3)	0.61560 (16)	0.0177 (3)	0.683 (3)
O1	0.0398 (3)	0.3931 (2)	0.34150 (16)	0.0214 (4)	
N1	0.3814 (3)	0.3582 (3)	0.19518 (18)	0.0165 (4)	
H1	0.3243	0.3831	0.1247	0.020*	
N2	0.6346 (3)	0.2730 (3)	0.32364 (17)	0.0155 (4)	
H2	0.7665	0.2340	0.3513	0.019*	
C1	0.6123 (4)	0.3176 (3)	0.1982 (2)	0.0161 (4)	
C2	0.2501 (4)	0.3552 (3)	0.3155 (2)	0.0165 (5)	
C3	0.4173 (3)	0.2981 (3)	0.4032 (2)	0.0153 (4)	
C4	0.3529 (4)	0.2791 (3)	0.5309 (2)	0.0172 (5)	
H4	0.1912	0.3066	0.5592	0.021*	
C5	0.484 (2)	0.225 (8)	0.6318 (12)	0.017 (2)	0.683 (3)
C6	0.388 (2)	0.234 (2)	0.7571 (9)	0.0201 (7)	0.683 (3)
H6	0.2288	0.2766	0.7856	0.024*	0.683 (3)
C7	0.5563 (7)	0.1717 (7)	0.8394 (5)	0.0218 (9)	0.683 (3)
H7	0.5228	0.1714	0.9280	0.026*	0.683 (3)
C8	0.7709 (9)	0.1127 (9)	0.7737 (4)	0.0227 (12)	0.683 (3)
H8	0.9045	0.0645	0.8122	0.027*	0.683 (3)
S2'	0.3819 (11)	0.2103 (10)	0.7847 (4)	0.0201 (7)	0.317
C5'	0.494 (4)	0.210 (17)	0.628 (2)	0.017 (2)	0.317
C6'	0.731 (3)	0.159 (3)	0.6117 (17)	0.0177 (3)	0.317
H6'	0.8277	0.1592	0.5304	0.021*	0.317 (3)
C7'	0.816 (3)	0.106 (2)	0.7307 (10)	0.0227 (12)	0.317
H7'	0.9731	0.0609	0.7379	0.027*	0.317 (3)
C8'	0.6415 (17)	0.1274 (17)	0.8312 (14)	0.0218 (9)	0.317
H8'	0.6628	0.0994	0.9177	0.026*	0.317 (3)

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
S1	0.0143 (3)	0.0310 (3)	0.0138 (3)	-0.0020 (2)	-0.0009 (2)	-0.0024 (2)
S2	0.0127 (7)	0.0198 (7)	0.0184 (5)	-0.0001 (5)	-0.0058 (5)	-0.0018 (4)
O1	0.0122 (7)	0.0315 (9)	0.0211 (9)	-0.0048 (6)	-0.0035 (6)	-0.0068 (7)

N1	0.0139 (8)	0.0223 (9)	0.0131 (9)	-0.0046 (7)	-0.0041 (7)	-0.0020 (7)
N2	0.0108 (8)	0.0213 (9)	0.0129 (9)	-0.0026 (6)	-0.0037 (7)	-0.0013 (7)
C1	0.0149 (10)	0.0173 (9)	0.0154 (11)	-0.0047 (7)	-0.0035 (8)	-0.0012 (8)
C2	0.0159 (10)	0.0186 (10)	0.0157 (11)	-0.0062 (8)	-0.0021 (8)	-0.0035 (8)
C3	0.0137 (10)	0.0162 (9)	0.0153 (11)	-0.0041 (7)	-0.0034 (8)	-0.0014 (8)
C4	0.0155 (10)	0.0182 (10)	0.0180 (11)	-0.0057 (8)	-0.0026 (8)	-0.0034 (8)
C5	0.0229 (15)	0.014 (7)	0.0159 (13)	-0.007 (2)	-0.0053 (11)	-0.0011 (14)
C6	0.0226 (7)	0.0262 (17)	0.011 (2)	-0.0063 (8)	-0.0009 (15)	-0.0042 (16)
C7	0.034 (3)	0.020 (2)	0.0125 (14)	-0.0076 (19)	-0.004 (2)	-0.0041 (14)
C8	0.031 (4)	0.0192 (13)	0.018 (3)	-0.0072 (17)	-0.012 (3)	0.003 (2)
S2'	0.0226 (7)	0.0262 (17)	0.011 (2)	-0.0063 (8)	-0.0009 (15)	-0.0042 (16)
C5'	0.0229 (15)	0.014 (7)	0.0159 (13)	-0.007 (2)	-0.0053 (11)	-0.0011 (14)
C6'	0.0127 (7)	0.0198 (7)	0.0184 (5)	-0.0001 (5)	-0.0058 (5)	-0.0018 (4)
C7'	0.031 (4)	0.0192 (13)	0.018 (3)	-0.0072 (17)	-0.012 (3)	0.003 (2)
C8'	0.034 (3)	0.020 (2)	0.0125 (14)	-0.0076 (19)	-0.004 (2)	-0.0041 (14)

Geometric parameters (Å, °)

S1—C1	1.660 (2)	C5—C6	1.380 (13)
S2—C5	1.706 (9)	C6—C7	1.433 (13)
S2—C8	1.702 (5)	C6—H6	0.9500
O1—C2	1.224 (3)	C7—C8	1.368 (5)
N1—C1	1.373 (3)	C7—H7	0.9500
N1—C2	1.374 (3)	C8—H8	0.9500
N1—H1	0.8800	S2'—C8'	1.693 (9)
N2—C1	1.358 (3)	S2'—C5'	1.705 (17)
N2—C3	1.407 (3)	C5'—C6'	1.378 (16)
N2—H2	0.8800	C6'—C7'	1.438 (15)
C2—C3	1.476 (3)	C6'—H6'	0.9500
C3—C4	1.344 (3)	C7'—C8'	1.356 (9)
C4—C5'	1.430 (8)	C7'—H7'	0.9500
C4—C5	1.431 (5)	C8'—H8'	0.9500
C4—H4	0.9500		
C5—S2—C8	92.3 (3)	C4—C5—S2	125.2 (7)
C1—N1—C2	111.81 (19)	C5—C6—C7	112.6 (10)
C1—N1—H1	124.1	C5—C6—H6	123.7
C2—N1—H1	124.1	C7—C6—H6	123.7
C1—N2—C3	110.50 (17)	C8—C7—C6	111.1 (7)
C1—N2—H2	124.8	C8—C7—H7	124.4
C3—N2—H2	124.8	C6—C7—H7	124.4
N2—C1—N1	107.33 (18)	C7—C8—S2	112.7 (5)
N2—C1—S1	126.45 (16)	C7—C8—H8	123.6
N1—C1—S1	126.20 (17)	S2—C8—H8	123.6
O1—C2—N1	126.4 (2)	C8'—S2'—C5'	93.5 (8)
O1—C2—C3	128.6 (2)	C6'—C5'—C4	128.4 (16)
N1—C2—C3	104.99 (17)	C6'—C5'—S2'	109.6 (10)
C4—C3—N2	132.2 (2)	C4—C5'—S2'	121.5 (14)

C4—C3—C2	122.56 (19)	C5'—C6'—C7'	113.2 (14)
N2—C3—C2	105.19 (18)	C5'—C6'—H6'	123.4
C3—C4—C5'	128.4 (9)	C7'—C6'—H6'	123.4
C3—C4—C5	131.6 (5)	C8'—C7'—C6'	111.3 (16)
C3—C4—H4	114.2	C8'—C7'—H7'	124.4
C5'—C4—H4	117.3	C6'—C7'—H7'	124.4
C5—C4—H4	114.2	C7'—C8'—S2'	112.2 (13)
C6—C5—C4	123.6 (9)	C7'—C8'—H8'	123.9
C6—C5—S2	111.2 (6)	S2'—C8'—H8'	123.9
C3—N2—C1—N1	-4.2 (2)	O1—C2—C3—C4	0.3 (3)
C3—N2—C1—S1	174.52 (16)	N1—C2—C3—C4	-179.96 (19)
C2—N1—C1—N2	4.3 (2)	O1—C2—C3—N2	-179.7 (2)
C2—N1—C1—S1	-174.39 (16)	N1—C2—C3—N2	0.1 (2)
C1—N1—C2—O1	177.1 (2)	C6—C7—C8—S2	1.0 (8)
C1—N1—C2—C3	-2.7 (2)	C5—S2—C8—C7	0.0 (18)
C1—N2—C3—C4	-177.4 (2)	C6'—C7'—C8'—S2'	-0.1 (18)
C1—N2—C3—C2	2.5 (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>
N2—H2...O1 ⁱ	0.88	2.20	2.873 (2)	133

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