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## Structure Reports

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# {3-[(2-Chloro-1,3-thiazol-4-yl)methyl]-1,3-thiazolidin-2-ylideneamino}-formonitrile

Ying-qi Li

Vocational–Technical Institute, Xiangtan University, Xiangtan 411100, People's Republic of China

Correspondence e-mail: liyingqi01@163.com

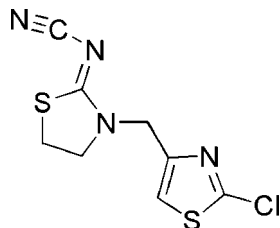
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Key indicators: single-crystal X-ray study;  $T = 294$  K; mean  $\sigma(\text{C}–\text{C}) = 0.003$  Å;  $R$  factor = 0.036;  $wR$  factor = 0.103; data-to-parameter ratio = 16.3.

In the title compound,  $\text{C}_8\text{H}_7\text{ClN}_4\text{S}_2$ , the dihedral angle between the thiazolidine ring (r.m.s. deviation = 0.028 Å) and the thiazole ring (r.m.s. deviation = 0.004 Å) is 74.74 (6)°. The formonitrile group is almost coplanar with the attached ring [C–N–C–N torsion angle = 167 (2)°].

## Related literature

For the biological activity of compounds containing a thiazole ring, see: Ehrenfreund *et al.* (2003); Kim *et al.* (2002); Maienfisch & Gsell (1998); Shiga *et al.* (2003); Smith & Hunter (2001); Tanaka *et al.* (2005). For the bioactivity of 1,3-thiazolidine derivatives, see: Albrecht *et al.* (2005); Liu & Li (2000); Ueda *et al.* (2004); Yeh & Chen (2002).



## Experimental

## Crystal data

$\text{C}_8\text{H}_7\text{ClN}_4\text{S}_2$	$V = 1088.5$ (3) Å <sup>3</sup>
$M_r = 258.75$	$Z = 4$
Monoclinic, $P2_1/n$	Mo $K\alpha$ radiation
$a = 6.1731$ (9) Å	$\mu = 0.70$ mm <sup>-1</sup>
$b = 16.807$ (2) Å	$T = 294$ K
$c = 10.9057$ (14) Å	$0.22 \times 0.20 \times 0.18$ mm
$\beta = 105.846$ (2)°	

## Data collection

Bruker SMART CCD area-detector diffractometer	6141 measured reflections
Absorption correction: multi-scan (SADABS; Sheldrick, 1996)	2220 independent reflections
$T_{\min} = 0.860$ , $T_{\max} = 0.884$	1862 reflections with $I > 2\sigma(I)$
	$R_{\text{int}} = 0.021$

## Refinement

$R[F^2 > 2\sigma(F^2)] = 0.036$	136 parameters
$wR(F^2) = 0.103$	H-atom parameters constrained
$S = 1.05$	$\Delta\rho_{\text{max}} = 0.30$ e Å <sup>-3</sup>
2220 reflections	$\Delta\rho_{\text{min}} = -0.34$ e Å <sup>-3</sup>

Data collection: SMART (Bruker, 2000); cell refinement: SAINT (Bruker, 2000); data reduction: SAINT; program(s) used to solve structure: SHELXS97 (Sheldrick, 2008); program(s) used to refine structure: SHELXL97 (Sheldrick, 2008); molecular graphics: SHELXTL (Sheldrick, 2008); software used to prepare material for publication: SHELXTL.

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: CI5173).

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## supporting information

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**{3-[(2-Chloro-1,3-thiazol-4-yl)methyl]-1,3-thiazolidin-2-ylideneamino}formo-nitrile****Ying-qi Li****S1. Comment**

Recently, compounds containing thiazole ring have been reported to possess various biological activities such as fungicidal, insecticidal, and anticancer activities (Maienfisch & Gsell, 1998; Smith *et al.*, 2001; Kim *et al.*, 2002; Ehrenfreund *et al.*, 2003; Shiga *et al.*, 2003; Tanaka *et al.*, 2005). In addition, 1,3-thiazolidine ring is an important heterocycle scaffold among thiazole compounds. In the past few years lots of 1,3-thiazolidine derivatives have attracted intense attention in medicinal research due to their broad spectrum bioactivities (Liu *et al.*, 2000; Yeh *et al.*, 2002; Ueda *et al.*, 2004; Albrecht *et al.*, 2005). In order to discover more biologically active thiazole compounds, we synthesized thiazole compounds containing 1,3-thiazolidine ring and we report here the crystal structure of the title compound.

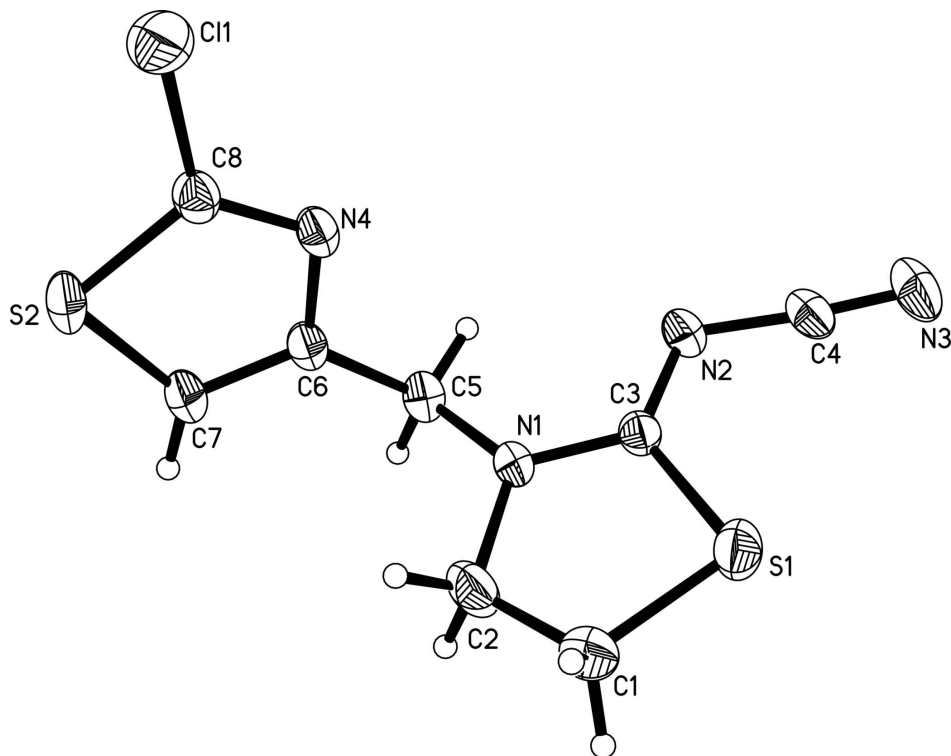
The molecule of the title compound (Fig.1) contains two planar rings, the substituted 1,3-thiazolidine ring (S1/C1/C2/N1/C3, r.m.s. deviation 0.028 Å) and the thiazole ring (S2/C8/N4/C6/C7, r.m.s. deviation 0.004 Å). The dihedral angle between the planes of 1,3-thiazolidine ring and thiazole ring is 74.74 (6)°.

**S2. Experimental**

To a stirred solution of 2-cyanoimino-1,3-thiazolidine (1.27 g, 0.01 mol), potassium carbonate (1.66 g, 0.012 mol) in 20 ml of acetonitrile was added dropwise a solution of 2-chloro-4-(chloromethyl)thiazole (1.68 g, 0.01 mol) in 15 ml of acetonitrile. The reaction mixture was heated to 333 K for 12 h and then filtered. The solvent was removed to give a solid product, which was recrystallized from ethyl acetate to afford colourless crystals.

**S3. Refinement**

All H atoms were placed in calculated positions, with C–H = 0.93 and 0.97 Å, and included in the final cycles of refinement using a riding model, with  $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$ .

**Figure 1**

The molecular structure of the title compound, showing the atom-numbering scheme. Displacement ellipsoids are drawn at the 30% probability level.

### {3-[(2-Chloro-1,3-thiazol-4-yl)methyl]-1,3-thiazolidin-2-ylideneamino}formonitrile

#### Crystal data

$C_8H_7ClN_4S_2$

$M_r = 258.75$

Monoclinic,  $P2_1/n$

Hall symbol:  $-P\ 2_1n$

$a = 6.1731\ (9)\ \text{\AA}$

$b = 16.807\ (2)\ \text{\AA}$

$c = 10.9057\ (14)\ \text{\AA}$

$\beta = 105.846\ (2)^\circ$

$V = 1088.5\ (3)\ \text{\AA}^3$

$Z = 4$

$F(000) = 528$

$D_x = 1.579\ \text{Mg m}^{-3}$

Mo  $K\alpha$  radiation,  $\lambda = 0.71073\ \text{\AA}$

Cell parameters from 3431 reflections

$\theta = 2.4\text{--}26.2^\circ$

$\mu = 0.70\ \text{mm}^{-1}$

$T = 294\ \text{K}$

Monoclinic, colourless

$0.22 \times 0.20 \times 0.18\ \text{mm}$

#### Data collection

Bruker SMART CCD area-detector  
diffractometer

Radiation source: fine-focus sealed tube

Graphite monochromator

$\varphi$  and  $\omega$  scans

Absorption correction: multi-scan

(*SADABS*; Sheldrick, 1996)

$T_{\min} = 0.860$ ,  $T_{\max} = 0.884$

6141 measured reflections

2220 independent reflections

1862 reflections with  $I > 2\sigma(I)$

$R_{\text{int}} = 0.021$

$\theta_{\max} = 26.4^\circ$ ,  $\theta_{\min} = 2.3^\circ$

$h = -7 \rightarrow 7$

$k = -15 \rightarrow 21$

$l = -11 \rightarrow 13$

*Refinement*Refinement on  $F^2$ 

Least-squares matrix: full

 $R[F^2 > 2\sigma(F^2)] = 0.036$  $wR(F^2) = 0.103$  $S = 1.05$ 

2220 reflections

136 parameters

0 restraints

Primary atom site location: structure-invariant  
direct methodsSecondary atom site location: difference Fourier  
mapHydrogen site location: inferred from  
neighbouring sites

H-atom parameters constrained

 $w = 1/[\sigma^2(F_o^2) + (0.0573P)^2 + 0.3256P]$ where  $P = (F_o^2 + 2F_c^2)/3$  $(\Delta/\sigma)_{\max} = 0.001$  $\Delta\rho_{\max} = 0.30 \text{ e } \text{\AA}^{-3}$  $\Delta\rho_{\min} = -0.34 \text{ e } \text{\AA}^{-3}$ *Special details*

**Geometry.** All e.s.d.'s (except the e.s.d. in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell e.s.d.'s are taken into account individually in the estimation of e.s.d.'s in distances, angles and torsion angles; correlations between e.s.d.'s in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell e.s.d.'s is used for estimating e.s.d.'s involving l.s. planes.

**Refinement.** Refinement of  $F^2$  against ALL reflections. The weighted  $R$ -factor  $wR$  and goodness of fit  $S$  are based on  $F^2$ , conventional  $R$ -factors  $R$  are based on  $F$ , with  $F$  set to zero for negative  $F^2$ . The threshold expression of  $F^2 > \sigma(F^2)$  is used only for calculating  $R$ -factors(gt) *etc.* and is not relevant to the choice of reflections for refinement.  $R$ -factors based on  $F^2$  are statistically about twice as large as those based on  $F$ , and  $R$ -factors based on ALL data will be even larger.

*Fractional atomic coordinates and isotropic or equivalent isotropic displacement parameters ( $\text{\AA}^2$ )*

	$x$	$y$	$z$	$U_{\text{iso}}^*/U_{\text{eq}}$
Cl1	-0.27073 (11)	0.02267 (4)	0.65605 (6)	0.0705 (2)
S1	0.70384 (10)	0.09234 (4)	1.21031 (5)	0.0588 (2)
S2	0.07788 (10)	0.10824 (3)	0.56567 (5)	0.05097 (18)
N1	0.4957 (3)	0.15672 (9)	0.99768 (14)	0.0389 (4)
N2	0.3327 (3)	0.18765 (11)	1.15862 (15)	0.0466 (4)
N3	0.3408 (4)	0.18457 (14)	1.38596 (19)	0.0669 (6)
N4	0.0452 (3)	0.11665 (10)	0.79535 (15)	0.0410 (4)
C1	0.8149 (4)	0.07222 (18)	1.0769 (2)	0.0660 (7)
H1A	0.9724	0.0873	1.0975	0.079*
H1B	0.8034	0.0159	1.0571	0.079*
C2	0.6843 (4)	0.11841 (17)	0.9667 (2)	0.0631 (7)
H2A	0.7801	0.1583	0.9440	0.076*
H2B	0.6296	0.0834	0.8942	0.076*
C3	0.4872 (3)	0.15132 (11)	1.11773 (17)	0.0370 (4)
C4	0.3448 (4)	0.18300 (13)	1.2819 (2)	0.0478 (5)
C5	0.3492 (3)	0.20877 (12)	0.90364 (17)	0.0426 (4)
H5A	0.4394	0.2499	0.8790	0.051*
H5B	0.2428	0.2346	0.9417	0.051*
C6	0.2233 (3)	0.16430 (11)	0.78796 (17)	0.0378 (4)
C7	0.2661 (4)	0.16624 (12)	0.67255 (17)	0.0451 (5)
H7	0.3818	0.1951	0.6541	0.054*
C8	-0.0433 (3)	0.08538 (11)	0.68491 (19)	0.0428 (4)

Atomic displacement parameters ( $\text{\AA}^2$ )

	$U^{11}$	$U^{22}$	$U^{33}$	$U^{12}$	$U^{13}$	$U^{23}$
Cl1	0.0659 (4)	0.0730 (4)	0.0743 (4)	-0.0187 (3)	0.0221 (3)	-0.0253 (3)
S1	0.0642 (4)	0.0707 (4)	0.0407 (3)	0.0175 (3)	0.0133 (3)	0.0086 (3)
S2	0.0674 (4)	0.0577 (3)	0.0290 (3)	0.0066 (3)	0.0152 (2)	-0.0030 (2)
N1	0.0391 (8)	0.0492 (9)	0.0307 (8)	0.0022 (7)	0.0131 (6)	-0.0022 (7)
N2	0.0483 (9)	0.0586 (10)	0.0382 (9)	0.0007 (8)	0.0208 (7)	-0.0049 (7)
N3	0.0869 (15)	0.0758 (14)	0.0499 (11)	-0.0106 (11)	0.0390 (11)	-0.0082 (10)
N4	0.0470 (9)	0.0462 (9)	0.0335 (8)	0.0025 (7)	0.0174 (7)	-0.0024 (7)
C1	0.0546 (13)	0.0878 (18)	0.0591 (14)	0.0197 (13)	0.0218 (11)	0.0030 (13)
C2	0.0664 (15)	0.0811 (17)	0.0517 (13)	0.0266 (13)	0.0328 (12)	0.0063 (12)
C3	0.0369 (9)	0.0420 (10)	0.0328 (9)	-0.0068 (8)	0.0108 (7)	-0.0030 (7)
C4	0.0526 (12)	0.0519 (12)	0.0465 (12)	-0.0061 (9)	0.0264 (9)	-0.0056 (9)
C5	0.0517 (11)	0.0417 (10)	0.0356 (9)	0.0012 (8)	0.0140 (8)	-0.0005 (8)
C6	0.0448 (10)	0.0398 (10)	0.0310 (9)	0.0074 (8)	0.0141 (7)	0.0027 (7)
C7	0.0549 (12)	0.0498 (11)	0.0345 (10)	0.0026 (9)	0.0191 (9)	0.0019 (8)
C8	0.0470 (10)	0.0435 (10)	0.0397 (10)	0.0051 (8)	0.0149 (8)	-0.0041 (8)

Geometric parameters ( $\text{\AA}$ ,  $^\circ$ )

Cl1—C8	1.715 (2)	N4—C6	1.380 (2)
S1—C3	1.7471 (19)	C1—C2	1.472 (3)
S1—C1	1.801 (2)	C1—H1A	0.97
S2—C7	1.709 (2)	C1—H1B	0.97
S2—C8	1.711 (2)	C2—H2A	0.97
N1—C3	1.328 (2)	C2—H2B	0.97
N1—C2	1.448 (3)	C5—C6	1.490 (3)
N1—C5	1.459 (2)	C5—H5A	0.97
N2—C3	1.309 (2)	C5—H5B	0.97
N2—C4	1.328 (3)	C6—C7	1.355 (3)
N3—C4	1.142 (3)	C7—H7	0.93
N4—C8	1.291 (2)		
C3—S1—C1	92.29 (10)	N2—C3—N1	122.18 (17)
C7—S2—C8	88.08 (10)	N2—C3—S1	125.57 (14)
C3—N1—C2	116.74 (17)	N1—C3—S1	112.25 (14)
C3—N1—C5	123.40 (16)	N3—C4—N2	173.6 (3)
C2—N1—C5	119.09 (15)	N1—C5—C6	111.98 (15)
C3—N2—C4	118.19 (18)	N1—C5—H5A	109.2
C8—N4—C6	108.81 (16)	C6—C5—H5A	109.2
C2—C1—S1	108.40 (16)	N1—C5—H5B	109.2
C2—C1—H1A	110.0	C6—C5—H5B	109.2
S1—C1—H1A	110.0	H5A—C5—H5B	107.9
C2—C1—H1B	110.0	C7—C6—N4	115.37 (17)
S1—C1—H1B	110.0	C7—C6—C5	125.85 (19)
H1A—C1—H1B	108.4	N4—C6—C5	118.78 (16)
N1—C2—C1	109.89 (18)	C6—C7—S2	110.57 (16)

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N1—C2—H2A	109.7	C6—C7—H7	124.7
C1—C2—H2A	109.7	S2—C7—H7	124.7
N1—C2—H2B	109.7	N4—C8—S2	117.15 (16)
C1—C2—H2B	109.7	N4—C8—C11	122.55 (16)
H2A—C2—H2B	108.2	S2—C8—C11	120.29 (12)
C3—S1—C1—C2	4.2 (2)	C3—N1—C5—C6	-124.90 (19)
C3—N1—C2—C1	6.9 (3)	C2—N1—C5—C6	65.5 (2)
C5—N1—C2—C1	177.2 (2)	C8—N4—C6—C7	-1.2 (2)
S1—C1—C2—N1	-6.7 (3)	C8—N4—C6—C5	178.42 (17)
C4—N2—C3—N1	-175.95 (18)	N1—C5—C6—C7	-106.0 (2)
C4—N2—C3—S1	3.3 (3)	N1—C5—C6—N4	74.4 (2)
C2—N1—C3—N2	175.7 (2)	N4—C6—C7—S2	0.9 (2)
C5—N1—C3—N2	5.9 (3)	C5—C6—C7—S2	-178.68 (15)
C2—N1—C3—S1	-3.7 (2)	C8—S2—C7—C6	-0.28 (16)
C5—N1—C3—S1	-173.46 (14)	C6—N4—C8—S2	1.0 (2)
C1—S1—C3—N2	-179.83 (19)	C6—N4—C8—C11	-179.65 (14)
C1—S1—C3—N1	-0.49 (17)	C7—S2—C8—N4	-0.42 (17)
C3—N2—C4—N3	167 (2)	C7—S2—C8—C11	-179.83 (14)

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