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(Z)-3-Allyl-5-(3-methoxybenzylidene)-2-sulfanyl- idene-1,3-thiazolidin-4-one

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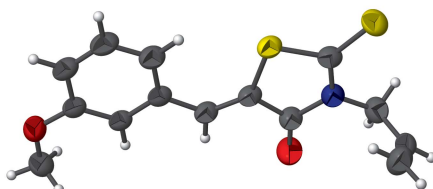
Keywords: crystal structure; rhodanine-based molecule; hydrogen bonds.

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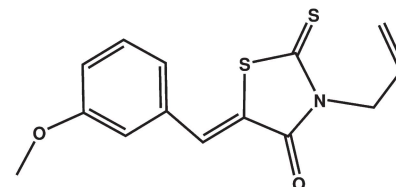
Structural data: full structural data are available from iucrdata.iucr.org

In the title compound, C₁₄H₁₃NO₂S₂, the rhodanine ring and the 3-methoxybenzylidene ring are nearly coplanar, as indicated by the dihedral angle of 1.77 (6)° between their planes. The allyl group is nearly perpendicular to the rhodanine ring, with a dihedral angle of 83.64 (19)°. An intramolecular C—H···S interaction forms an S(6) ring motif. In the crystal, molecules are linked by pairs of C—H···O hydrogen bonds into inversion dimers.

3D view



Chemical scheme



Structure description

Compounds containing 2-thioxothiazolidin-4-one (rhodanine) and its derivatives have been reported to exhibit a broad spectrum of biological activities, acting as antidiabetic, anticancer, antitubercular, anti-HIV and antiparasitic agents (Murugan *et al.*, 2009; Chandrappa *et al.*, 2009; Mallikarjuna *et al.*, 2009; Murugesan *et al.*, 2011; Zhang *et al.*, 2009). The unusual biological activity displayed by many rhodanine-based molecules has made them attractive synthetic targets.

The molecule of the title compound is built up from a rhodanine ring (S1/N1/C8–C10) linked to an allyl group (C11–C13) at the nitrogen atom and to a 3-methoxybenzylidene ring (C1–C6) as shown in Fig. 1. In the crystal, molecules are linked by pairs of C—H···O hydrogen bonds (Table 1), forming an inversion dimer as shown in Fig. 2.

Synthesis and crystallization

To a solution of 3-allylrhodanine (1.15 mmol, 0.2 g) in 10 ml of THF, (3-methoxybenzylidene)-4-methyl-5-oxopyrazolidin-2-ium-1-ide (1.38 mmol) was added. The

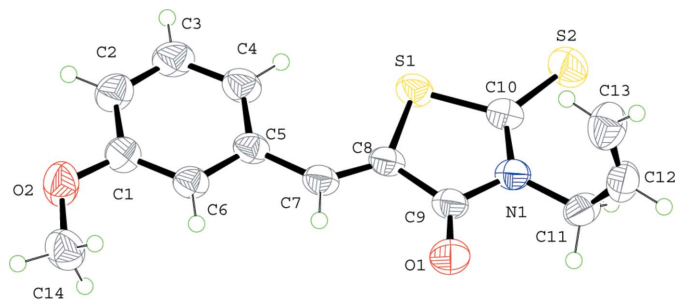


Figure 1

The molecular structure of the title compound with the atom-labelling scheme. Displacement ellipsoids are drawn at the 50% probability level. H atoms are represented as small circles.

mixture was refluxed for 8 h until the reaction was completed (TLC) and a yellow spot (TLC $R_f = 0.3$, using hexane/ethyl acetate 1:9) was generated cleanly. The solvent was evaporated *in vacuo*. The crude product was purified on silica gel using hexane/ethyl acetate (1:9) as eluent. The title compound was recrystallized from ethanol (yield 78%, m.p. 364 K).

Refinement

Crystal data, data collection and structure refinement details are summarized in Table 2. The reflection (0 0 1) was affected by the beam-stop and was removed during refinement.

Acknowledgements

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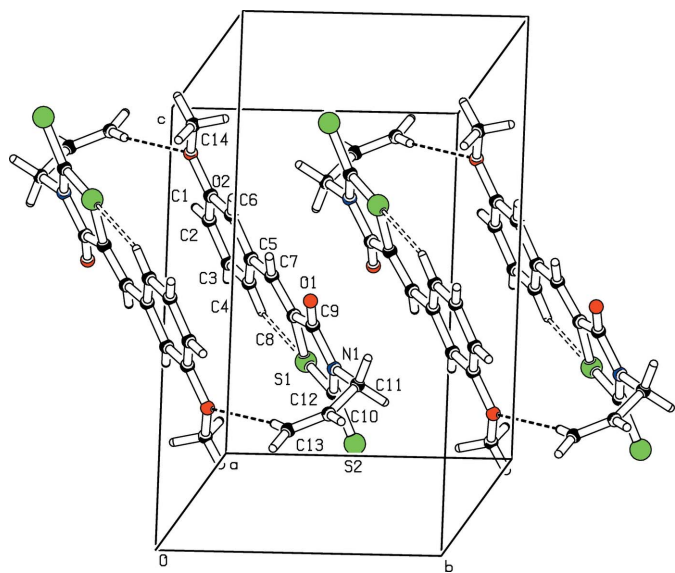


Figure 2

A crystal packing diagram of the title compound, showing the hydrogen bonds as dashed lines.

Table 1

Hydrogen-bond geometry (\AA , $^\circ$).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
$C4-H4\cdots S1$	0.93	2.55	3.2497 (17)	133
$C13-H13A\cdots O2^i$	0.93	2.57	3.441 (2)	157

Symmetry code: (i) $-x + 1, -y + 2, -z + 1$.

Table 2

Experimental details.

Crystal data	
Chemical formula	$C_{14}H_{13}NO_2S_2$
M_r	291.37
Crystal system, space group	Triclinic, $P\bar{1}$
Temperature (K)	296
a, b, c (\AA)	6.9841 (14), 8.3241 (18), 13.116 (3)
α, β, γ ($^\circ$)	89.276 (9), 75.614 (9), 72.095 (10)
V (\AA^3)	701.2 (3)
Z	2
Radiation type	Mo $K\alpha$
μ (mm^{-1})	0.38
Crystal size (mm)	$0.31 \times 0.27 \times 0.21$
Data collection	
Diffractometer	Bruker X8 APEX diffractometer
Absorption correction	Multi-scan (SADABS; Bruker, 2009)
T_{\min}, T_{\max}	0.479, 0.746
No. of measured, independent and observed [$I > 2\sigma(I)$] reflections	28065, 4522, 3618
R_{int}	0.029
$(\sin \theta/\lambda)_{\text{max}}$ (\AA^{-1})	0.729
Refinement	
$R[F^2 > 2\sigma(F^2)], wR(F^2), S$	0.039, 0.117, 1.02
No. of reflections	4522
No. of parameters	172
H-atom treatment	H-atom parameters constrained
$\Delta\rho_{\text{max}}, \Delta\rho_{\text{min}}$ ($e \text{\AA}^{-3}$)	0.30, -0.26

Computer programs: APEX2 (Bruker, 2009), SAINT (Bruker, 2009), SHELXS2014 (Sheldrick, 2008), SHELXL2014 (Sheldrick, 2015), ORTEP-3 for Windows (Farrugia, 2012), publCIF (Westrip, 2010).

References

- Bruker (2009). APEX2, SAINT and SADABS. Bruker AXS Inc., Madison, Wisconsin, USA.
- Chandrappa, S., Kavitha, C. V., Shahabuddin, M. S., Vinaya, K., Ananda Kumar, C. S., Ranganatha, S. R., Raghavan, S. C. & Rangappa, K. S. (2009). *Bioorg. Med. Chem.* **17**, 2576–2584.
- Farrugia, L. J. (2012). *J. Appl. Cryst.* **45**, 849–854.
- Mallikarjuna, B. P., Sastry, B. S., Suresh Kumar, G. V., Rajendraprasad, Y., Chandrashekar, S. M. & Sathisha, K. (2009). *Eur. J. Med. Chem.* **44**, 4739–4746.
- Murugan, R., Anbazhagan, S. & Sriman Narayanan, S. (2009). *Eur. J. Med. Chem.* **44**, 3272–3279.
- Murugesan, V., Tiwari, V. S., Saxena, R., Tripathi, R., Paranjape, R., Kulkarni, S., Makwana, N., Suryawanshi, R. & Katti, S. B. (2011). *Bioorg. Med. Chem.* **19**, 6919–6926.
- Sheldrick, G. M. (2008). *Acta Cryst.* **A64**, 112–122.
- Sheldrick, G. M. (2015). *Acta Cryst.* **C71**, 3–8.
- Westrip, S. P. (2010). *J. Appl. Cryst.* **43**, 920–925.
- Zhang, X., Li, X., Li, D., Qu, G., Wang, J., Loiseau, P. M. & Fan, X. (2009). *Bioorg. Med. Chem. Lett.* **19**, 6280–6283.

full crystallographic data

IUCrData (2016). **1**, x160052 [doi:10.1107/S2414314616000523]

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(Z)-3-Allyl-5-(3-methoxybenzylidene)-2-sulfanylidene-1,3-thiazolidin-4-one*Crystal data*

$C_{14}H_{13}NO_2S_2$

$M_r = 291.37$

Triclinic, $P\bar{1}$

$a = 6.9841$ (14) Å

$b = 8.3241$ (18) Å

$c = 13.116$ (3) Å

$\alpha = 89.276$ (9)°

$\beta = 75.614$ (9)°

$\gamma = 72.095$ (10)°

$V = 701.2$ (3) Å³

$Z = 2$

$F(000) = 304$

$D_x = 1.380$ Mg m⁻³

Melting point: 364 K

Mo $K\alpha$ radiation, $\lambda = 0.71073$ Å

Cell parameters from 4522 reflections

$\theta = 2.6$ – 31.2 °

$\mu = 0.38$ mm⁻¹

$T = 296$ K

Block, colourless

$0.31 \times 0.27 \times 0.21$ mm

Data collection

Bruker X8 APEX
diffractometer

Radiation source: fine-focus sealed tube

Graphite monochromator

φ and ω scans

Absorption correction: multi-scan
(*SADABS*; Bruker, 2009)

$T_{\min} = 0.479$, $T_{\max} = 0.746$

28065 measured reflections

4522 independent reflections

3618 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.029$

$\theta_{\max} = 31.2$ °, $\theta_{\min} = 2.6$ °

$h = -10 \rightarrow 10$

$k = -12 \rightarrow 12$

$l = -19 \rightarrow 19$

Refinement

Refinement on F^2

Least-squares matrix: full

$R[F^2 > 2\sigma(F^2)] = 0.039$

$wR(F^2) = 0.117$

$S = 1.02$

4522 reflections

172 parameters

0 restraints

Hydrogen site location: inferred from
neighbouring sites

H-atom parameters constrained

$w = 1/[\sigma^2(F_o^2) + (0.0624P)^2 + 0.1273P]$

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

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

$\Delta\rho_{\max} = 0.30$ e Å⁻³

$\Delta\rho_{\min} = -0.26$ e Å⁻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.

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

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$
C1	0.6784 (2)	0.92899 (16)	0.26792 (11)	0.0453 (3)
C2	0.8814 (2)	0.88347 (19)	0.27837 (12)	0.0528 (3)
H2	0.9846	0.9093	0.2274	0.063*
C3	0.9283 (2)	0.8002 (2)	0.36440 (12)	0.0555 (3)
H3	1.0641	0.7700	0.3711	0.067*
C4	0.7776 (2)	0.76011 (19)	0.44141 (11)	0.0512 (3)
H4	0.8122	0.7042	0.4993	0.061*
C5	0.57287 (19)	0.80427 (16)	0.43169 (10)	0.0410 (2)
C6	0.52494 (19)	0.88990 (16)	0.34378 (10)	0.0418 (3)
H6	0.3895	0.9205	0.3365	0.050*
C7	0.40267 (19)	0.76903 (16)	0.50824 (10)	0.0431 (3)
H7	0.2737	0.8106	0.4932	0.052*
C8	0.40122 (19)	0.68638 (16)	0.59698 (10)	0.0408 (2)
C9	0.2049 (2)	0.66565 (17)	0.66206 (10)	0.0450 (3)
C10	0.4403 (2)	0.52824 (16)	0.76057 (10)	0.0434 (3)
C11	0.0714 (2)	0.53236 (19)	0.82335 (12)	0.0520 (3)
H11A	0.1286	0.4223	0.8486	0.062*
H11B	-0.0273	0.5227	0.7851	0.062*
C12	-0.0422 (2)	0.65574 (19)	0.91596 (12)	0.0533 (3)
H12	-0.1612	0.6390	0.9590	0.064*
C13	0.0068 (3)	0.7842 (2)	0.94329 (14)	0.0677 (4)
H13A	0.1244	0.8067	0.9030	0.081*
H13B	-0.0757	0.8538	1.0031	0.081*
C14	0.4469 (3)	1.0541 (2)	0.16012 (13)	0.0616 (4)
H14A	0.4481	1.1117	0.0962	0.092*
H14B	0.4124	0.9525	0.1529	0.092*
H14C	0.3452	1.1265	0.2178	0.092*
N1	0.24062 (17)	0.57615 (14)	0.75017 (9)	0.0435 (2)
O1	0.03559 (17)	0.71622 (17)	0.64424 (9)	0.0660 (3)
O2	0.64706 (17)	1.01213 (16)	0.18002 (9)	0.0643 (3)
S1	0.60384 (5)	0.59150 (4)	0.65517 (3)	0.04671 (11)
S2	0.52699 (7)	0.42768 (6)	0.85588 (3)	0.06213 (13)

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
C1	0.0380 (6)	0.0469 (6)	0.0453 (6)	-0.0068 (5)	-0.0083 (5)	-0.0009 (5)
C2	0.0346 (6)	0.0607 (8)	0.0560 (8)	-0.0105 (5)	-0.0042 (5)	-0.0015 (6)
C3	0.0317 (6)	0.0695 (9)	0.0612 (9)	-0.0085 (6)	-0.0135 (6)	-0.0020 (7)
C4	0.0391 (6)	0.0636 (8)	0.0488 (7)	-0.0090 (6)	-0.0162 (5)	0.0004 (6)
C5	0.0352 (5)	0.0458 (6)	0.0395 (6)	-0.0076 (4)	-0.0110 (4)	-0.0062 (5)
C6	0.0332 (5)	0.0470 (6)	0.0422 (6)	-0.0077 (4)	-0.0100 (4)	-0.0035 (5)
C7	0.0354 (5)	0.0534 (7)	0.0409 (6)	-0.0113 (5)	-0.0135 (5)	-0.0028 (5)
C8	0.0349 (5)	0.0488 (6)	0.0392 (6)	-0.0105 (5)	-0.0132 (4)	-0.0048 (5)
C9	0.0406 (6)	0.0549 (7)	0.0441 (6)	-0.0171 (5)	-0.0166 (5)	0.0017 (5)

C10	0.0416 (6)	0.0445 (6)	0.0449 (6)	-0.0107 (5)	-0.0161 (5)	-0.0023 (5)
C11	0.0508 (7)	0.0571 (8)	0.0593 (8)	-0.0288 (6)	-0.0195 (6)	0.0116 (6)
C12	0.0428 (7)	0.0607 (8)	0.0539 (8)	-0.0159 (6)	-0.0095 (6)	0.0197 (6)
C13	0.0682 (10)	0.0646 (9)	0.0590 (9)	-0.0200 (8)	0.0030 (8)	-0.0032 (7)
C14	0.0578 (9)	0.0692 (9)	0.0569 (9)	-0.0132 (7)	-0.0219 (7)	0.0129 (7)
N1	0.0412 (5)	0.0495 (6)	0.0440 (5)	-0.0169 (4)	-0.0150 (4)	0.0016 (4)
O1	0.0423 (5)	0.0990 (9)	0.0671 (7)	-0.0278 (5)	-0.0264 (5)	0.0233 (6)
O2	0.0462 (6)	0.0820 (8)	0.0585 (6)	-0.0147 (5)	-0.0103 (5)	0.0227 (6)
S1	0.03460 (15)	0.0597 (2)	0.04389 (18)	-0.00911 (13)	-0.01372 (12)	0.00062 (13)
S2	0.0570 (2)	0.0726 (3)	0.0594 (2)	-0.01587 (19)	-0.02600 (18)	0.01825 (18)

Geometric parameters (Å, °)

C1—O2	1.3668 (17)	C9—N1	1.4001 (17)
C1—C6	1.3860 (18)	C10—N1	1.3685 (17)
C1—C2	1.3920 (19)	C10—S2	1.6335 (14)
C2—C3	1.373 (2)	C10—S1	1.7494 (14)
C2—H2	0.9300	C11—N1	1.4639 (17)
C3—C4	1.385 (2)	C11—C12	1.487 (2)
C3—H3	0.9300	C11—H11A	0.9700
C4—C5	1.4008 (18)	C11—H11B	0.9700
C4—H4	0.9300	C12—C13	1.298 (2)
C5—C6	1.4050 (18)	C12—H12	0.9300
C5—C7	1.4564 (18)	C13—H13A	0.9300
C6—H6	0.9300	C13—H13B	0.9300
C7—C8	1.3441 (19)	C14—O2	1.422 (2)
C7—H7	0.9300	C14—H14A	0.9600
C8—C9	1.4827 (18)	C14—H14B	0.9600
C8—S1	1.7445 (13)	C14—H14C	0.9600
C9—O1	1.2079 (16)		
O2—C1—C6	124.67 (12)	N1—C10—S2	127.52 (11)
O2—C1—C2	115.21 (13)	N1—C10—S1	110.74 (10)
C6—C1—C2	120.12 (13)	S2—C10—S1	121.74 (8)
C3—C2—C1	119.57 (13)	N1—C11—C12	114.59 (11)
C3—C2—H2	120.2	N1—C11—H11A	108.6
C1—C2—H2	120.2	C12—C11—H11A	108.6
C2—C3—C4	121.39 (13)	N1—C11—H11B	108.6
C2—C3—H3	119.3	C12—C11—H11B	108.6
C4—C3—H3	119.3	H11A—C11—H11B	107.6
C3—C4—C5	119.69 (14)	C13—C12—C11	127.06 (14)
C3—C4—H4	120.2	C13—C12—H12	116.5
C5—C4—H4	120.2	C11—C12—H12	116.5
C4—C5—C6	118.87 (12)	C12—C13—H13A	120.0
C4—C5—C7	124.13 (12)	C12—C13—H13B	120.0
C6—C5—C7	117.00 (11)	H13A—C13—H13B	120.0
C1—C6—C5	120.35 (12)	O2—C14—H14A	109.5
C1—C6—H6	119.8	O2—C14—H14B	109.5

C5—C6—H6	119.8	H14A—C14—H14B	109.5
C8—C7—C5	130.67 (12)	O2—C14—H14C	109.5
C8—C7—H7	114.7	H14A—C14—H14C	109.5
C5—C7—H7	114.7	H14B—C14—H14C	109.5
C7—C8—C9	120.39 (11)	C10—N1—C9	116.48 (11)
C7—C8—S1	130.08 (10)	C10—N1—C11	123.16 (12)
C9—C8—S1	109.53 (9)	C9—N1—C11	120.32 (11)
O1—C9—N1	123.01 (13)	C1—O2—C14	118.75 (12)
O1—C9—C8	126.62 (13)	C8—S1—C10	92.87 (6)
N1—C9—C8	110.37 (11)		

Hydrogen-bond geometry (Å, °)

<i>D</i> —H \cdots <i>A</i>	<i>D</i> —H	H \cdots <i>A</i>	<i>D</i> \cdots <i>A</i>	<i>D</i> —H \cdots <i>A</i>
C4—H4 \cdots S1	0.93	2.55	3.2497 (17)	133
C13—H13 <i>A</i> \cdots O2 ⁱ	0.93	2.57	3.441 (2)	157

Symmetry code: (i) $-x+1, -y+2, -z+1$.