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 Methyl 3-[(*E,E*)-3-phenylprop-2-enylidene]dithiocarbazate

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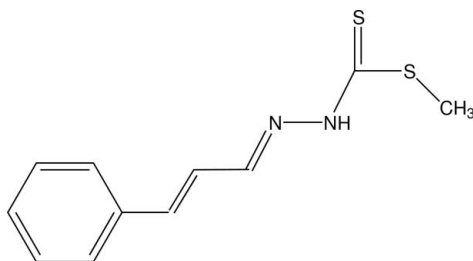
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 Key indicators: single-crystal X-ray study; $T = 293$ K; mean $\sigma(\text{C}-\text{C}) = 0.004$ Å; R factor = 0.038; wR factor = 0.116; data-to-parameter ratio = 14.8.

In the title compound, $\text{C}_{11}\text{H}_{12}\text{N}_2\text{S}_2$, the dithiocarbazate group adopts an *EE* configuration with respect to the $\text{C}=\text{C}$ and $\text{C}=\text{N}$ bonds of the propenylidene group. The atoms of the propenylidene and dithiocarbazate unit are essentially coplanar, with a maximum deviation of 0.058 (1) Å; the phenyl ring forms a dihedral angle of 18.3 (1)° with this fragment. In the crystal, molecules form inversion dimers *via* pairs of $\text{N}-\text{H}\cdots\text{S}$ hydrogen bonds involving the terminal S atom.

Related literature

 For the synthesis and a related structure, see: Tarafder *et al.* (2008).


Experimental

Crystal data

 $\text{C}_{11}\text{H}_{12}\text{N}_2\text{S}_2$
 $M_r = 236.35$

 Monoclinic, $P2_1/c$
 $a = 10.408$ (2) Å
 $b = 5.4950$ (9) Å
 $c = 20.988$ (2) Å
 $\beta = 100.697$ (10)°
 $V = 1179.5$ (3) Å³
 $Z = 4$
 Mo $K\alpha$ radiation
 $\mu = 0.42$ mm⁻¹
 $T = 293$ K
 $0.40 \times 0.15 \times 0.12$ mm

Data collection

 Enraf–Nonius DIP1030 image-plate diffractometer
 6316 measured reflections

 2048 independent reflections
 1569 reflections with $I > 2.0\sigma(I)$
 $R_{\text{int}} = 0.037$

Refinement

 $R[F^2 > 2\sigma(F^2)] = 0.038$
 $wR(F^2) = 0.116$
 $S = 1.04$
 2048 reflections

 138 parameters
 H-atom parameters constrained
 $\Delta\rho_{\text{max}} = 0.14$ e Å⁻³
 $\Delta\rho_{\text{min}} = -0.21$ e Å⁻³

Table 1

Hydrogen-bond geometry (Å, °).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
$\text{N1}-\text{H1}\cdots\text{S1}^i$	0.86	2.67	3.4086 (19)	145

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

Data collection: *XPRESS* (MacScience, 2002); cell refinement: *DENZO* (Otwinowski & Minor, 1997); data reduction: *DENZO* and *SCALEPACK* (Otwinowski & Minor, 1997); program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *ORTEP-3 for Windows* (Farrugia, 1997); software used to prepare material for publication: *WinGX* (Farrugia, 1999).

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: PV2336).

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Methyl 3-[(*E,E*)-3-phenylprop-2-enylidene]dithiocarbazate

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S1. Comment

The molecule of the title compound is shown in Fig. 1. The atoms of propenylidene and dithiocarbazate moiety lie essentially in the same plane, with a maximum deviation from the mean plane of 0.058 (1) Å exhibited by S2, while the phenyl ring forms a dihedral angle of 18.3 (1)° with the cited fragment. The crystal packing evidences molecules connected in pairs about inversion centers and connected by N1—H1⋯S1 hydrogen bonds involving the terminal sulfur atom (Table 1). The structural features of the title compound are similar to those observed in a benzyl derivative (Tarafder *et al.*, 2008). The Schiff base is potentially bidentate and coordinates *via* the β-nitrogen and the thiolate anion generated during complexation.

S2. Experimental

The *S*-methylthiocarbazate (2.44 g, 0.2 mol), prepared as previously described (Tarafder *et al.*, 2008), was dissolved in hot absolute ethanol (30–40 ml). To this solution an equimolar amount of cinnamaldehyde in hot absolute ethanol (20 ml) was added and the mixture was heated for 20 min and then cooled. The orange precipitate thus formed was separated and dried *in vacuo* over anhydrous CaCl₂. Orange needle shaped single crystals of the Schiff base were obtained after recrystallization from acetone over 15 days; *M. p.* 443 K (very sharp and abrupt).

S3. Refinement

All H atoms were located geometrically and treated as riding atoms, with N—H = 0.86 and C—H = 0.93 and 0.96 Å for aryl and methyl H-atoms with $U_{iso}(H) = 1.2U_{eq}(N \text{ or aryl-C})$ or $1.5U_{eq}(\text{methyl-C})$.

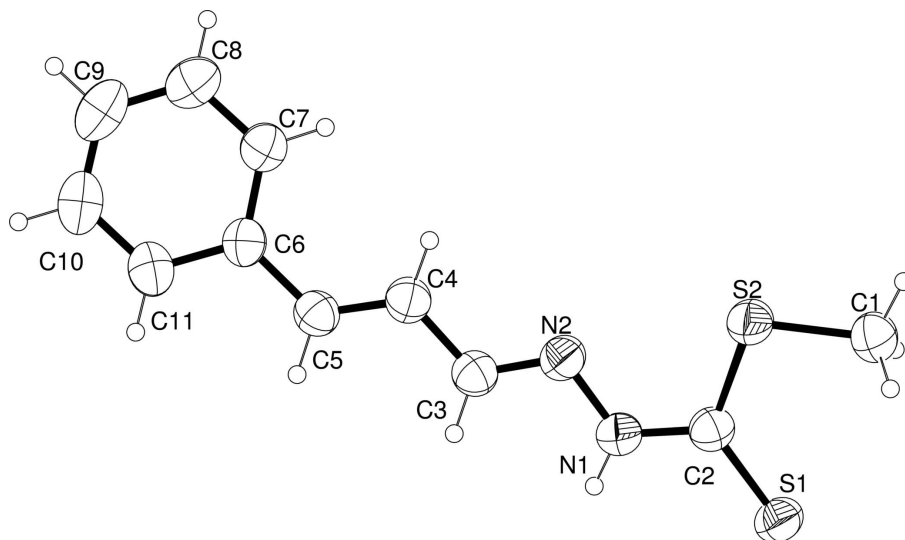


Figure 1

ORTEP drawing of the title molecule; thermal ellipsoids are drawn at the 40% probability level.

Methyl 3-[(*E,E*)-3-phenylprop-2-enylidene]dithiocarbazate

Crystal data

$C_{11}H_{12}N_2S_2$

$M_r = 236.35$

Monoclinic, $P2_1/c$

Hall symbol: $-P\ 2ybc$

$a = 10.408\ (2)\ \text{\AA}$

$b = 5.4950\ (9)\ \text{\AA}$

$c = 20.988\ (2)\ \text{\AA}$

$\beta = 100.697\ (10)^\circ$

$V = 1179.5\ (3)\ \text{\AA}^3$

$Z = 4$

$F(000) = 496$

$D_x = 1.331\ \text{Mg m}^{-3}$

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

Cell parameters from 275 reflections

$\theta = 3.6\text{--}20.7^\circ$

$\mu = 0.42\ \text{mm}^{-1}$

$T = 293\ \text{K}$

Needle, orange

$0.40 \times 0.15 \times 0.12\ \text{mm}$

Data collection

Enraf–Nonius DIP1030 image-plate
diffractometer

Radiation source: fine-focus sealed tube

Graphite monochromator

φ -scans with narrow frames

6316 measured reflections

2048 independent reflections

1569 reflections with $I > 2.0\sigma(I)$

$R_{\text{int}} = 0.037$

$\theta_{\text{max}} = 25.3^\circ$, $\theta_{\text{min}} = 3.1^\circ$

$h = -12 \rightarrow 12$

$k = -6 \rightarrow 6$

$l = -25 \rightarrow 25$

Refinement

Refinement on F^2

Least-squares matrix: full

$R[F^2 > 2\sigma(F^2)] = 0.038$

$wR(F^2) = 0.116$

$S = 1.04$

2048 reflections

138 parameters

0 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

$w = 1/[\sigma^2(F_o^2) + (0.0721P)^2]$

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

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

$\Delta\rho_{\text{max}} = 0.14\ \text{e \AA}^{-3}$

$\Delta\rho_{\text{min}} = -0.21\ \text{e \AA}^{-3}$

Extinction correction: *SHELXL*,
 $F_c^* = kF_c [1 + 0.001 \times F_c^2 \lambda^3 / \sin(2\theta)]^{-1/4}$

Extinction coefficient: 0.023 (4)

Special details

Geometry. All s.u.'s (except the s.u. in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell s.u.'s are taken into account individually in the estimation of s.u.'s in distances, angles and torsion angles; correlations between s.u.'s in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell s.u.'s is used for estimating s.u.'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 $F^2 > 2\sigma(F^2)$ 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 (\AA^2)

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$
S1	1.15322 (6)	0.68232 (10)	0.57196 (3)	0.0759 (2)
S2	1.08034 (6)	0.51686 (10)	0.69852 (3)	0.0734 (2)
N1	0.99960 (18)	0.3187 (3)	0.58644 (8)	0.0682 (5)
H1	0.9962	0.2915	0.5458	0.082*
N2	0.92717 (19)	0.1751 (3)	0.62061 (9)	0.0680 (5)
C1	1.1813 (3)	0.7789 (4)	0.72034 (11)	0.0791 (6)
H1A	1.2653	0.7525	0.7089	0.119*
H1B	1.1918	0.8059	0.7662	0.119*
H1C	1.1407	0.9187	0.6976	0.119*
C2	1.0751 (2)	0.4996 (3)	0.61523 (10)	0.0625 (5)
C3	0.8633 (2)	0.0018 (3)	0.58766 (11)	0.0656 (5)
H3	0.8723	-0.0225	0.5449	0.079*
C4	0.7788 (2)	-0.1538 (3)	0.61562 (10)	0.0670 (5)
H4	0.7718	-0.1294	0.6587	0.080*
C5	0.7095 (2)	-0.3328 (3)	0.58259 (10)	0.0653 (5)
H5	0.7240	-0.3613	0.5408	0.078*
C6	0.6139 (2)	-0.4880 (3)	0.60503 (10)	0.0634 (5)
C7	0.5705 (3)	-0.4487 (4)	0.66304 (12)	0.0750 (6)
H7	0.6053	-0.3208	0.6898	0.090*
C8	0.4770 (3)	-0.5962 (5)	0.68147 (14)	0.0898 (7)
H8	0.4487	-0.5659	0.7202	0.108*
C9	0.4250 (3)	-0.7874 (5)	0.64316 (17)	0.0963 (9)
H9	0.3623	-0.8870	0.6560	0.116*
C10	0.4657 (3)	-0.8309 (4)	0.58615 (16)	0.0944 (8)
H10	0.4305	-0.9603	0.5601	0.113*
C11	0.5596 (3)	-0.6831 (4)	0.56675 (13)	0.0799 (7)
H11	0.5866	-0.7146	0.5277	0.096*

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
S1	0.0761 (5)	0.0872 (4)	0.0662 (4)	-0.0152 (3)	0.0178 (3)	0.0061 (2)
S2	0.0809 (5)	0.0799 (4)	0.0592 (4)	-0.0143 (3)	0.0122 (3)	0.0035 (2)

N1	0.0743 (13)	0.0726 (10)	0.0600 (10)	-0.0089 (9)	0.0181 (9)	-0.0002 (7)
N2	0.0681 (12)	0.0698 (10)	0.0670 (10)	-0.0066 (8)	0.0150 (9)	0.0030 (8)
C1	0.0811 (17)	0.0813 (13)	0.0730 (14)	-0.0110 (12)	0.0092 (12)	-0.0046 (10)
C2	0.0574 (13)	0.0664 (12)	0.0636 (12)	0.0012 (9)	0.0112 (9)	0.0024 (8)
C3	0.0686 (15)	0.0647 (12)	0.0631 (12)	0.0011 (10)	0.0111 (10)	0.0014 (8)
C4	0.0698 (15)	0.0673 (11)	0.0631 (12)	-0.0028 (10)	0.0104 (10)	0.0010 (9)
C5	0.0682 (14)	0.0658 (12)	0.0611 (11)	0.0013 (10)	0.0099 (10)	-0.0002 (8)
C6	0.0633 (14)	0.0560 (10)	0.0683 (13)	0.0023 (8)	0.0057 (10)	0.0036 (8)
C7	0.0799 (17)	0.0700 (12)	0.0756 (14)	-0.0059 (11)	0.0160 (12)	0.0006 (10)
C8	0.090 (2)	0.0870 (15)	0.0964 (18)	-0.0020 (14)	0.0276 (15)	0.0160 (13)
C9	0.0772 (18)	0.0813 (16)	0.129 (3)	-0.0085 (13)	0.0151 (17)	0.0274 (16)
C10	0.091 (2)	0.0722 (15)	0.114 (2)	-0.0154 (13)	0.0022 (17)	-0.0017 (13)
C11	0.0866 (18)	0.0680 (13)	0.0813 (15)	-0.0038 (12)	0.0061 (13)	-0.0054 (10)

Geometric parameters (Å, °)

S1—C2	1.664 (2)	C5—C6	1.454 (3)
S2—C2	1.741 (2)	C5—H5	0.9300
S2—C1	1.791 (2)	C6—C7	1.392 (3)
N1—C2	1.339 (3)	C6—C11	1.395 (3)
N1—N2	1.381 (2)	C7—C8	1.376 (3)
N1—H1	0.8600	C7—H7	0.9300
N2—C3	1.287 (3)	C8—C9	1.372 (4)
C1—H1A	0.9600	C8—H8	0.9300
C1—H1B	0.9600	C9—C10	1.363 (4)
C1—H1C	0.9600	C9—H9	0.9300
C3—C4	1.428 (3)	C10—C11	1.388 (4)
C3—H3	0.9300	C10—H10	0.9300
C4—C5	1.335 (3)	C11—H11	0.9300
C4—H4	0.9300		
C2—S2—C1	102.02 (11)	C4—C5—H5	116.6
C2—N1—N2	121.39 (18)	C6—C5—H5	116.6
C2—N1—H1	119.3	C7—C6—C11	117.4 (2)
N2—N1—H1	119.3	C7—C6—C5	123.05 (18)
C3—N2—N1	114.92 (18)	C11—C6—C5	119.5 (2)
S2—C1—H1A	109.5	C8—C7—C6	121.1 (2)
S2—C1—H1B	109.5	C8—C7—H7	119.5
H1A—C1—H1B	109.5	C6—C7—H7	119.5
S2—C1—H1C	109.5	C9—C8—C7	120.6 (3)
H1A—C1—H1C	109.5	C9—C8—H8	119.7
H1B—C1—H1C	109.5	C7—C8—H8	119.7
N1—C2—S1	120.44 (16)	C10—C9—C8	119.7 (3)
N1—C2—S2	113.58 (15)	C10—C9—H9	120.1
S1—C2—S2	125.98 (12)	C8—C9—H9	120.1
N2—C3—C4	121.2 (2)	C9—C10—C11	120.4 (2)
N2—C3—H3	119.4	C9—C10—H10	119.8
C4—C3—H3	119.4	C11—C10—H10	119.8

C5—C4—C3	122.8 (2)	C10—C11—C6	120.8 (3)
C5—C4—H4	118.6	C10—C11—H11	119.6
C3—C4—H4	118.6	C6—C11—H11	119.6
C4—C5—C6	126.9 (2)		
C2—N1—N2—C3	-177.27 (18)	C4—C5—C6—C11	-173.6 (2)
N2—N1—C2—S1	-175.70 (15)	C11—C6—C7—C8	-0.5 (3)
N2—N1—C2—S2	4.4 (2)	C5—C6—C7—C8	178.1 (2)
C1—S2—C2—N1	-177.67 (16)	C6—C7—C8—C9	0.6 (4)
C1—S2—C2—S1	2.41 (18)	C7—C8—C9—C10	-0.4 (4)
N1—N2—C3—C4	-176.76 (18)	C8—C9—C10—C11	0.1 (4)
N2—C3—C4—C5	178.9 (2)	C9—C10—C11—C6	0.0 (4)
C3—C4—C5—C6	-174.93 (19)	C7—C6—C11—C10	0.2 (3)
C4—C5—C6—C7	7.9 (3)	C5—C6—C11—C10	-178.4 (2)

Hydrogen-bond geometry (Å, °)

<i>D—H...A</i>	<i>D—H</i>	<i>H...A</i>	<i>D...A</i>	<i>D—H...A</i>
N1—H1...S1 ⁱ	0.86	2.67	3.4086 (19)	145

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