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Key indicators

Single-crystal X-ray study T = 293 KMean σ (C–C) = 0.005 Å R factor = 0.051 wR factor = 0.132 Data-to-parameter ratio = 13.5

For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e. In the title compound, $C_{12}H_{13}N_3OS$, the molecular backbone formed by the thiazolidinone and 2-(propan-2-ylidene)hydrazone groups is planar. There are some weak intra- and intermolecular hydrogen-bond interactions which stabilize the crystal structure.

(Z)-3-Phenyl-2-(propan-2-ylidenehydrazono)-

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Comment

thiazolidin-4-one

It has been reported that both thiazoles and thiazolidines have good and wide insecticidal, fungicidal, herbicidal and acaricidal activity (Melnikov, *et al.*, 1979; Kratt & Salberk, 1984). As part of our search for compounds with good herbicidal and fungicidal activity, the title compound, (I), was synthesized. We report here its crystal structure.



Bond lengths and angles in the thiazolidine ring (Table 1) are in agreement those reported previously (Buyuktimkin et al., 1984; Yang et al., 1995; Pomés Hernandez et al., 1996). The molecular backbone formed by the thiazolidinone and 2-(propan-2-ylidene)hydrazone groups is essentially planar, the maximum deviations being -0.070 (3) and 0.029 (3)° for C2 and N2, respectively. The phenyl ring is twisted by $70.98 (7)^{\circ}$ with respect to this plane. There are some weak $C-H \cdots O$ and C-H···N intra- and intermolecular hydrogen-bond interactions which stabilize the crystal structure (Table 2).

Experimental

A mixture of hydrazine (0.02 mol), ethyl chloroacetate (0.02 mol)and phenyl isothiocyanate (0.02 mol) was stirred in refluxing acetone (5 ml) for 5 h at 327 K to afford the title compound (3.71 g, yield75%). Single crystals suitable for X-ray measurements were obtained by recrystallization from ethanol at room temperature.

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organic papers

Crystal data

C12H13N3OS $M_r = 247.31$ Monoclinic, $P2_1/c$ a = 16.859(9) Åb = 5.667 (3) Åc = 12.580(7) Å $\beta = 90.685 (9)^{\circ}$ $V = 1201.8 (11) \text{ Å}^3$ Z = 4

Data collection

Bruker SMART CCD area-detector diffractometer φ and ω scans Absorption correction: multi-scan (SADABS; Sheldrick, 1996) $T_{\min} = 0.950, T_{\max} = 0.970$ 6139 measured reflections

Refinement

Refinement on F^2	$w = 1/[\sigma^2(F_o^2) + (0$
$R[F^2 > 2\sigma(F^2)] = 0.051$	where $P = (F_o^2 +$
$wR(F^2) = 0.132$	$(\Delta/\sigma)_{\rm max} = 0.001$
S = 0.89	$\Delta \rho_{\rm max} = 0.27 \ {\rm e} \ {\rm \AA}^{-2}$
2121 reflections	$\Delta \rho_{\rm min} = -0.28 \text{ e Å}$
157 parameters	Extinction correction
H-atom parameters constrained	Extinction coefficie

Table 1

Selected geometric parameters (Å, °).

S1-C1	1.739 (3)	N2-C1	1.277 (4)
S1-C2	1.790 (3)	N2-N3	1.418 (4)
N1-C3	1.386 (4)	N3-C10	1.261 (4)
N1-C1	1.387 (4)		
C1-S1-C2	91.87 (15)	C1-N2-N3	107.7 (3)
C3-N1-C1	116.0 (3)	N2-C1-S1	125.3 (2)
C3-N1-C4	119.5 (2)	N1-C1-S1	112.2 (2)
C1-N1-C4	124.5 (3)		

Table 2

Hydrogen-bonding geometry (Å, °).

$D - H \cdot \cdot \cdot A$	D-H	$H \cdot \cdot \cdot A$	$D \cdot \cdot \cdot A$	$D - \mathbf{H} \cdot \cdot \cdot A$
$\begin{array}{c} \hline C12-H12A\cdots N2\\ C5-H5\cdots O1^{i}\\ C8-H8\cdots O1^{ii} \end{array}$	0.96	2.30	2.739 (5)	107
	0.93	2.41	3.321 (4)	165
	0.93	2.50	3.413 (4)	168

Symmetry codes: (i) x, y - 1, z; (ii) $x, \frac{3}{2} - y, \frac{1}{2} + z$.

 $D_x = 1.367 \text{ Mg m}^{-3}$ Mo $K\alpha$ radiation Cell parameters from 1528 reflections $\theta=2.4{-}22.4^\circ$ $\mu = 0.26 \text{ mm}^{-3}$ T = 293 (2) KBlock, yellow $0.18 \times 0.16 \times 0.12 \text{ mm}$

2121 independent reflections 1093 reflections with $I > 2\sigma(I)$ $R_{\rm int} = 0.075$ $\theta_{\rm max} = 25.0^{\circ}$ $h = -20 \rightarrow 11$ $k = -6 \rightarrow 6$ $l = -14 \rightarrow 14$

 $(.0657P)^2$ $+2F_c^2)/3$ -3 on: SHELXL97 ent: 0.010 (2)





View of (I), with displacement ellipsoids drawn at the 40% probability level.

All H atoms were placed in calculated positions, with C-H = 0.93-0.97 Å, and refined using a riding model, with $U_{iso}(H) = 1.2U_{eq}(C)$ for the aryl and methylene H atoms and $1.5U_{eq}(C)$ for the methyl H atoms.

Data collection: SMART (Bruker, 1998); cell refinement: SAINT (Bruker, 1999); data reduction: SAINT; program(s) used to solve structure: SHELXS97 (Sheldrick, 1997); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: SHELXTL (Bruker, 1999); software used to prepare material for publication: SHELXTL.

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