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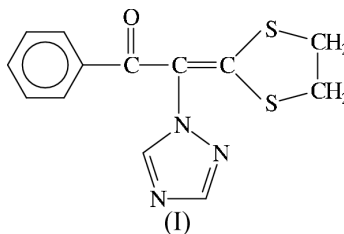
## Key indicators

Single-crystal X-ray study  
 $T = 293$  K  
Mean  $\sigma(\text{C}-\text{C}) = 0.005$  Å  
 $R$  factor = 0.047  
 $wR$  factor = 0.103  
Data-to-parameter ratio = 15.4For details of how these key indicators were  
automatically derived from the article, see  
<http://journals.iucr.org/e>.2-(1,3-Dithiolan-2-ylidene)-1-phenyl-2-  
(1,2,4-triazol-1-yl)ethanone

In the title compound,  $\text{C}_{13}\text{H}_{11}\text{N}_3\text{OS}_2$ , the dihedral angles between the planes of the triazole and phenyl rings and that through the four  $\text{NC}_3$  atoms of the  $\text{C}=\text{C}$  are  $76. (1)$  and  $57.6 (1)^\circ$ , respectively. There are some intermolecular interactions in the crystal structure.

## Comment

Many *N*-heterocyclic compounds have been widely used as potent and broad-spectrum fungicides, such as buthilbate, fenarimol and nuarimol (Kato *et al.*, 1975; Gestel *et al.*, 1980). Compounds containing the triazole ring system are well known as efficient fungicides in agriculture and medicine, where they act by inhibiting the biosynthesis of ergosterol (Banting *et al.*, 1989). They have also good plant-growth regulatory activity on a wide variety of crops (Xu *et al.*, 2002). In order to search for new triazole compounds with higher bioactivity, the title compound, (I), was synthesized and its structure is reported here.



In compound (I), the bond lengths and angles are generally normal in the phenyl and triazole rings (Ji *et al.*, 2002). The  $\text{C}8-\text{C}9$  bond length of  $1.370 (4)$  Å is indicative of considerable double-bond character. The six atoms  $\text{S}1$ ,  $\text{S}2$ ,  $\text{N}3$ ,  $\text{C}7$ ,  $\text{C}8$  and  $\text{C}9$  lie in a plane ( $P1$ ). The dihedral angles formed by the triazole ring with the phenyl ring ( $\text{C}1/\text{C}2/\text{C}3/\text{C}4/\text{C}5/\text{C}6$ ) and  $P1$  are  $79.1 (1)$  and  $76.5 (1)^\circ$ , respectively. The dihedral angle between the phenyl ring and  $P1$  is  $57.6 (1)^\circ$ . Torsion angles are given in Table 1.

The conformation of the dithiacyclopentane ring is half-chair, with a total puckering amplitude (Cremer & Pople, 1975)  $Q_T = 0.456 (3)$  Å and a pseudo-twofold axis running along the direction through  $\text{C}9$  and the mid-point of the  $\text{C}10-\text{C}11$  bond.

There are some weak intermolecular  $\text{C}-\text{H}\cdots\text{O}$  and  $\text{C}-\text{H}\cdots\text{N}$  hydrogen bonds (Table 2), which stabilize the structure of (I).

## Experimental

The title compound was prepared by reaction of  $\alpha$ -(1,2,4-triazol-1-yl) acetophenone (5.7 g, 0.02 mol),  $\text{CS}_2$  (1.9 g, 0.025 mol) and

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BrCH<sub>2</sub>CH<sub>2</sub>Br (0.025 mol) in a diethyl ether solution (40 ml) at room temperature. Single crystals of (I) suitable for X-ray measurements were obtained by recrystallization from chloroform–ethyl acetate (*v/v*, 1:3) at room temperature.

#### Crystal data

C<sub>13</sub>H<sub>11</sub>N<sub>3</sub>OS<sub>2</sub>  
*M<sub>r</sub>* = 289.37  
 Monoclinic, *P*2<sub>1</sub>/*c*  
*a* = 8.0523 (16) Å  
*b* = 10.170 (2) Å  
*c* = 17.116 (5) Å  
 $\beta$  = 112.63 (3)°  
*V* = 1293.8 (6) Å<sup>3</sup>  
*Z* = 4

*D<sub>x</sub>* = 1.486 Mg m<sup>-3</sup>  
 Mo *K*α radiation  
 Cell parameters from 20 reflections  
 $\theta$  = 2–11°  
 $\mu$  = 0.41 mm<sup>-1</sup>  
*T* = 293 (2) K  
 Block, yellow  
 0.20 × 0.20 × 0.18 mm

#### Data collection

Rigaku R-AXIS IV imaging-plate diffractometer  
 $\varphi$  scans  
 Absorption correction: none  
 4447 measured reflections  
 2642 independent reflections

1868 reflections with *I* > 2σ(*I*)  
*R*<sub>int</sub> = 0.036  
 $\theta_{\max}$  = 26.5°  
*h* = 0 → 10  
*k* = -13 → 12  
*l* = -22 → 20

#### Refinement

Refinement on *F*<sup>2</sup>  
*R*[*F*<sup>2</sup> > 2σ(*F*<sup>2</sup>)] = 0.047  
*wR*(*F*<sup>2</sup>) = 0.103  
*S* = 1.09  
 2642 reflections  
 172 parameters  
 H-atom parameters constrained

$w = 1/[\sigma^2(F_o^2) + (0.0338P)^2 + 0.668P]$   
 where  $P = (F_o^2 + 2F_c^2)/3$   
 $(\Delta/\sigma)_{\max} < 0.001$   
 $\Delta\rho_{\max} = 0.24 \text{ e \AA}^{-3}$   
 $\Delta\rho_{\min} = -0.40 \text{ e \AA}^{-3}$

**Table 1**

Selected torsion angles (°).

C5–C6–C7–C8	47.1 (5)	N2–N3–C8–C9	–98.4 (3)
N2–N3–C8–C7	78.5 (4)	S2–C9–C8–C7	–179.6 (2)
S1–C9–C8–C7	3.2 (5)	C1–C6–C7–O1	41.3 (4)

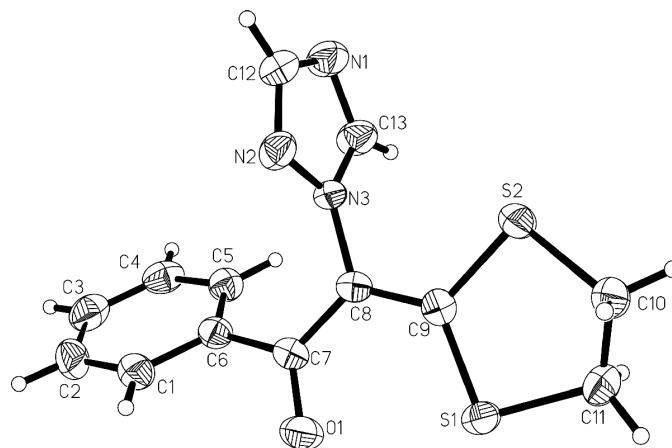
**Table 2**

Hydrogen-bonding 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>
C10–H10B···O1 <sup>i</sup>	0.97	2.58	3.422 (4)	146
C11–H11B···N2 <sup>i</sup>	0.97	2.60	3.559 (4)	170
C13–H13A···N2 <sup>ii</sup>	0.93	2.49	3.359 (4)	156

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

The H atoms were positioned geometrically and were treated as riding on their parent C atoms, with C–H distances in the range 0.93–0.97 Å and *U*<sub>iso</sub>(H) = 1.2*U*<sub>eq</sub>(C).



**Figure 1**

The structure of (I), showing 50% probability displacement ellipsoids and the atom-numbering scheme.

Data collection: *R-AXIS Software* (Rigaku, 1997); cell refinement: *R-AXIS Software*; data reduction: *R-AXIS Software*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *SHELXTL/PC* (Sheldrick, 1990); software used to prepare material for publication: *WinGX* (Farrugia, 1999).

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