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## Structure Reports

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## Key indicators

Single-crystal X-ray study
$T=294 \mathrm{~K}$
Mean $\sigma(\mathrm{C}-\mathrm{C})=0.005 \AA$
$R$ factor $=0.056$
$w R$ factor $=0.117$
Data-to-parameter ratio $=14.2$
For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e.

[^0]
## 2-Fluoro-5-methylphenyl (1H-1,2,4-triazol-1-yl)-(1,1,3-trioxo-1,2-dihydro-1,2-benzothiazol-2-yl)methyl ketone

In the title compound, $\mathrm{C}_{18} \mathrm{H}_{13} \mathrm{FN}_{4} \mathrm{O}_{4} \mathrm{~S}$, the dihedral angles between the planes of the triazole and the sulfobenzoimide ring and benzene ring are 87.69 (3) and $74.35(2)^{\circ}$, respectively. There are weak intermolecular hydrogen-bond interactions that assemble the molecules into slabs parallel to the $b c$ plane.

## Comment

As an important type of fungicide, triazole compounds are highly efficient and of low toxicity (Shi et al., 1995; Xu et al., 2002). On the other hand, sulfobenzoimide is used for diuretic (Chien, 1959), ataratic (Vnederlands, 1965) and soporific applications (Graadums et al., 1973), and some compounds containing the sulfobenzoimide group possess highly fungicidal and phytocidal activity (Hlasta et al., 1993; Hanagan \& Pasteris, 1986). Thus, since triazole compounds have a different fungicidal mechanism from that of sulfobenzoimide compounds, it was expected that a compound with both groups, such as title compound, (I) (Fig. 1), would have an enhanced fungicidal activity.

(I)

In (I), the $\mathrm{C} 9-\mathrm{N} 4$ and $\mathrm{C} 10-\mathrm{N} 3$ bond lengths (Table 1) are close to the typical $\mathrm{C}=\mathrm{N}$ double-bond length [1.330 (3) $\AA$; Dean, 1998), which reveals that these two bonds have features of unsaturated double bonds. The dihedral angles between the planes of the triazole and the sulfobenzoimide ring and benzene ring are 87.69 (3) and 74.35 (2) $)^{\circ}$, respectively. There are some weak $\mathrm{C}-\mathrm{H} \cdots \mathrm{O}$ and $\mathrm{C}-\mathrm{H} \cdots \mathrm{F}$ intermolecular hydrogen bonds in the crystal structure, assembling the molecules into slabs parallel to the $b c$ plane (Fig. 2, Table 2).

## Experimental

Triethylamine ( 0.022 mol ) was dissolved in acetone ( 5 ml ) and a mixture of 2-fluoro-2-(1H-1,2,4-triazole-1-yl)acetophenone $(0.02 \mathrm{~mol})$, sulfobenzoimide $(0.02 \mathrm{~mol})$ and acetone $(20 \mathrm{ml})$ was
$\qquad$
added dropwise with stirring at 273 K for 2 h , and then for 10 h at room temperature. The solution was then filtered, concentrated and purified by chromatography on a silica gel column to give a white solid (yield $34 \%$ ). Single crystals suitable for X-ray measurements were obtained by recrystallization from ethanol and ethyl acetate (1:2) at room temperature.

## Crystal data

$\mathrm{C}_{18} \mathrm{H}_{13} \mathrm{FN}_{4} \mathrm{O}_{4} \mathrm{~S}$
$M_{r}=400.38$
Monoclinic, C2/c
$a=21.269$ (5) A
$b=10.594$ (3) $\AA$
$c=15.758$ (4) $\AA$
$\beta=94.823$ (4) ${ }^{\circ}$
$V=3538.0(15) \AA^{3}$

## $Z=8$

$D_{x}=1.503 \mathrm{Mg} \mathrm{m}^{-3}$
Mo $K \alpha$ radiation
$\mu=0.23 \mathrm{~mm}^{-1}$
$T=294$ (2) K
Block, white
$0.22 \times 0.14 \times 0.06 \mathrm{~mm}$

## Data collection

Bruker SMART CCD area-detector diffractometer
$\varphi$ and $\omega$ scans
Absorption correction: multi-scan (SADABS; Sheldrick, 1996)

$$
T_{\min }=0.952, T_{\max }=0.987
$$

## Refinement

Refinement on $F^{2}$

$$
\begin{gathered}
w=1 /\left[\sigma^{2}\left(F_{\mathrm{o}}^{2}\right)+(0.0309 P)^{2}\right. \\
\quad+4.2871 P] \\
\text { where } P=\left(F_{\mathrm{o}}^{2}+2 F_{\mathrm{c}}^{2}\right) / 3 \\
(\Delta / \sigma)_{\max }<0.001 \\
\Delta \rho_{\max }=0.37 \mathrm{e}^{-3} \AA^{-3} \\
\Delta \rho_{\min }=-0.46 \mathrm{e}^{-3}
\end{gathered}
$$

9516 measured reflections 3609 independent reflections
2137 reflections with $I>2 \sigma(I)$
$R_{\text {int }}=0.060$
$\theta_{\text {max }}=26.4^{\circ}$
$w R\left(F^{2}\right)=0.117$
$S=1.07$
3609 reflections
254 parameters
H -atom parameters constrained

Table 1
Selected bond lengths ( $\AA$ ).

| F1-C13 | $1.358(4)$ | N4-C9 | $1.312(4)$ |
| :--- | :--- | :--- | :--- |
| N3-C10 | $1.322(4)$ |  |  |

Table 2
Hydrogen-bond geometry ( $\AA,{ }^{\circ}$ ).

| $D-\mathrm{H} \cdots A$ | $D-\mathrm{H}$ | $\mathrm{H} \cdots A$ | $D \cdots A$ | $D-\mathrm{H} \cdots A$ |
| :---: | :---: | :---: | :---: | :---: |
| $\mathrm{C} 5-\mathrm{H} 5 \cdots \mathrm{O} 4^{\text {i }}$ | 0.93 | 2.54 | 3.464 (4) | 173 |
| $\mathrm{C} 8-\mathrm{H} 8 \cdots \mathrm{O}^{\text {ii }}$ | 0.98 | 2.27 | 3.000 (4) | 130 |
| $\mathrm{C} 10-\mathrm{H} 10 \cdots \mathrm{O} 3{ }^{\text {iii }}$ | 0.93 | 2.45 | 3.375 (4) | 175 |
| C14-H14 $\ldots$ F1 ${ }^{\text {iv }}$ | 0.93 | 2.49 | 3.394 (4) | 163 |

Symmetry codes: (i) $-x+\frac{1}{2}, y-\frac{1}{2},-z+\frac{1}{2}$; (ii) $-x+\frac{1}{2}, y+\frac{1}{2},-z+\frac{1}{2}$; (iii) $x,-y+2, z+\frac{1}{2}$, (iv) $-x+\frac{1}{2},-y+\frac{5}{2},-z$.

All H atoms were placed in calculated positions, with $\mathrm{C}-\mathrm{H}=0.93-$ $0.98 \AA$, and refined using a riding model, with $U_{\text {iso }}(\mathrm{H})=1.2 U_{\text {eq }}(\mathrm{C})$, or $U_{\text {iso }}(\mathrm{H})=1.5 U_{\text {eq }}(\mathrm{C})$ for 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.


Figure 1
The molecular structure of (I), with displacement ellipsoids drawn at the 40\% probability level.


Figure 2
A packing diagram of the title compound; viewed down the $b$ axis. Hydrogen bonds are shown as dashed lines.

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