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## Structure Reports

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Key indicators
Single-crystal X-ray study
$T=298 \mathrm{~K}$
Mean $\sigma(\mathrm{C}-\mathrm{C})=0.004 \AA$
$R$ factor $=0.062$
$w R$ factor $=0.137$
Data-to-parameter ratio $=13.4$
For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e.

## 3-[6-(4-Methoxyphenyl)-7H-1,2,4-triazolo-[3,4-b][1,3,4]thiadiazin-3-yl]propan-1-ol

In the title compound, $\mathrm{C}_{14} \mathrm{H}_{16} \mathrm{~N}_{4} \mathrm{O}_{2} \mathrm{~S}$, the six-membered thiadiazine ring adopts a distorted boat conformation. $\mathrm{O}-$ $\mathrm{H} \cdots \mathrm{N}$ hydrogen bonds link the molecules into centrosymmetric dimers and enhance the stability of the crystal structure.

## Comment

As potentially biologically active reagents, 3,6-disubstituted$7 H-1,2,4$-triaozlo[3,4-b][1,3,4]thiadiazines have received considerable attention over the past two decades (Zhou et al., 2006; Nadkarni et al., 2001). Triazoles fused with thiadiazines have been shown to exhibit antimicrobial (Feng et al., 1992) and diuretic properties (Mohan \& Anjaneyulu, 1987) and to act as photographic couplers (Holla et al., 2001). In this paper, we report the synthesis and crystal structure of the title compound, (I).


In (I) (Fig. 1 and Table 1), the five-membered triazole ring is conjugated. The six-membered thiadiazine ring adopts a distorted boat conformation. $\mathrm{O}-\mathrm{H} \cdots \mathrm{N}$ hydrogen bonds (Table 2) link the molecules into pairs around a center of symmetry (Fig. 2), enhancing the stability of the crystal structure.

## Experimental

Carbon disulfide ( 13 ml ) and hydrazine hydrate ( 24 ml ) mixed with water ( 75 ml ) were refluxed for 1 h at 363 K to form thiocarbohydrazide. 1,4-Butyrolactone ( 0.01 mol ) and thiocarbohydrazide ( 0.01 mol ) were refluxed in pyridine $(40 \mathrm{ml})$ for 4 h to obtain 4 -amino-5-mercapto-3-(3-hydroxypropyl)-1,2,4-triazole, (II), following the method of Xiong et al. (2002). To a solution of (II) ( 0.01 mol ) in absolute ethanol ( 20 ml ), was added 2-bromo-1-(4-methoxyphenyl)ethanone ( 0.01 mol ). The mixture was refluxed for 7 h . The solid

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[^0]obtained on cooling was filtered, washed with cold water, dried and recrystallized from ethanol to give (I). The purified product was dissolved in $95 \%$ ethanol and single crystals were obtained after 4 d .

## Crystal data

$\mathrm{C}_{14} \mathrm{H}_{16} \mathrm{~N}_{4} \mathrm{O}_{2} \mathrm{~S}$
$M_{r}=304.38$
Monoclinic, $P 2_{1} / c$
$a=7.6166$ (7) A
$b=12.8582(12) \AA$
$c=15.9198$ (13) $\AA$
$\beta=111.228(4)^{\circ}$
$V=1453.3$ (2) $\AA^{3}$

## Data collection

Bruker APEX area-detector diffractometer
$\varphi$ and $\omega$ scans
Absorption correction: multi-scan (SADABS; Bruker, 2002)
$T_{\text {min }}=0.944, T_{\text {max }}=0.959$

## $Z=4$

$D_{x}=1.391 \mathrm{Mg} \mathrm{m}^{-3}$
Mo $K \alpha$ radiation $\mu=0.23 \mathrm{~mm}^{-1}$
$T=298$ (2) K
Block, colorless
$0.25 \times 0.17 \times 0.16 \mathrm{~mm}$

7493 measured reflections 2556 independent reflections 2288 reflections with $I>2 \sigma(I)$
$R_{\text {int }}=0.031$
$\theta_{\text {max }}=25.0^{\circ}$

## Refinement

Refinement on $F^{2}$
$R\left[F^{2}>2 \sigma\left(F^{2}\right)\right]=0.062$
$w R\left(F^{2}\right)=0.137$
$S=1.20$
2556 reflections
191 parameters
H -atom parameters constrained

$$
\begin{aligned}
& w=1 /\left[\sigma^{2}\left(F_{\mathrm{o}}^{2}\right)+(0.0504 P)^{2}\right. \\
& \quad+0.734 P] \\
& \quad \text { where } P=\left(F_{\mathrm{o}}^{2}+2 F_{\mathrm{c}}^{2}\right) / 3 \\
& (\Delta / \sigma)_{\max }=0.001 \\
& \Delta \rho_{\max }=0.24 \mathrm{e} \AA^{-3} \\
& \Delta \rho_{\min }=-0.31 \mathrm{e}^{-3}
\end{aligned}
$$

Table 1
Selected geometric parameters ( $\left(\AA^{\circ}{ }^{\circ}\right)$.

| S1-C10 | $1.731(3)$ | $\mathrm{N} 2-\mathrm{C} 10$ | $1.367(3)$ |
| :--- | :---: | :--- | :--- |
| S1-C9 | $1.808(3)$ | $\mathrm{N} 3-\mathrm{C} 10$ | $1.297(3)$ |
| N1-C8 | $1.283(3)$ | $\mathrm{N} 3-\mathrm{N} 4$ | $1.403(3)$ |
| $\mathrm{N} 1-\mathrm{N} 2$ | $1.393(3)$ | $\mathrm{N} 4-\mathrm{C} 11$ | $1.299(3)$ |
| N2-C11 | $1.365(3)$ | $\mathrm{C} 8-\mathrm{C} 9$ | $1.510(4)$ |
|  |  |  |  |
| $\mathrm{C} 10-\mathrm{S} 1-\mathrm{C} 9$ | $95.36(13)$ | $\mathrm{C} 5-\mathrm{C} 8-\mathrm{C} 9$ | $118.9(2)$ |
| $\mathrm{C} 8-\mathrm{N} 1-\mathrm{N} 2$ | $116.2(2)$ | $\mathrm{C} 8-\mathrm{C} 9-\mathrm{S} 1$ | $113.99(19)$ |
| $\mathrm{C} 11-\mathrm{N} 2-\mathrm{C} 10$ | $105.4(2)$ | $\mathrm{N} 3-\mathrm{C} 10-\mathrm{N} 2$ | $110.7(2)$ |
| C11-N2-N1 | $124.2(2)$ | $\mathrm{N} 3-\mathrm{C} 10-\mathrm{S} 1$ | $128.8(2)$ |
| C10-N2-N1 | $129.8(2)$ | $\mathrm{N} 2-\mathrm{C} 10-\mathrm{S} 1$ | $120.42(19)$ |
| C10-N3-N4 | $106.3(2)$ | $\mathrm{N} 4-\mathrm{C} 11-\mathrm{N} 2$ | $109.4(2)$ |
| C11-N4-N3 | $108.2(2)$ | $\mathrm{N} 4-\mathrm{C} 11-\mathrm{C} 12$ | $127.0(2)$ |
| N1-C8-C5 | $116.6(2)$ | $\mathrm{N} 2-\mathrm{C} 11-\mathrm{C} 12$ | $123.6(2)$ |
| N1-C8-C9 | $124.3(2)$ |  |  |

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

| $D-\mathrm{H} \cdots A$ | $D-\mathrm{H}$ | $\mathrm{H} \cdots A$ | $D \cdots A$ | $D-\mathrm{H} \cdots A$ |
| :--- | :--- | :--- | :--- | :--- |
| $\mathrm{O} 2-\mathrm{H} 2 \cdots \mathrm{~N} 4^{\mathrm{i}}$ | 0.82 | 2.08 | $2.892(3)$ | 173 |

Symmetry code: (i) $-x+2,-y+1,-z+1$.
All H atoms were positioned geometrically and allowed to ride on their parent atoms at distances of $\mathrm{Cs} p^{2}-\mathrm{H}=0.93 \AA$ with $U_{\text {iso }}=1.2 U_{\text {eq }}$ (parent atom), Csp $p^{3}-\mathrm{H}=0.97 \AA$ with $U_{\text {iso }}=1.5 U_{\text {eq }}$ (parent atom) and $\mathrm{O}-\mathrm{H}=0.82 \AA$ with $U_{\text {iso }}=1.5 U_{\text {eq }}$ (parent atom).

Data collection: SMART (Bruker, 2002); cell refinement: SAINT (Bruker, 2002); data reduction: SAINT; program(s) used to solve structure: SHELXS97 (Sheldrick, 1997); program(s) used to refine


Figure 1
The molecular structure of (I) with the atom numbering, showing displacement ellipsoids at the $30 \%$ probability level.


Figure 2
Packing diagram for (I), showing the hydrogen-bonded (dashed lines) dimers.
structure: SHELXL97 (Sheldrick, 1997); molecular graphics: SHELXTL (Bruker, 2002); software used to prepare material for publication: SHELXL97.

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