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Key indicators

Single-crystal X-ray study T = 273 K Mean σ (C–C) = 0.003 Å R factor = 0.038 wR factor = 0.115 Data-to-parameter ratio = 14.7

For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e. In the title compound, $C_7H_{12}N_4OS$, the triazine ring is planar. The crystal structure is stabilized by intermolecular $N-H\cdots O$ hydrogen bonds which link the molecules into chains running along the *c* axis.

1,2,4-triazin-5(2H)-one

4-Amino-6-tert-butyl-3-thioxo-3,4-dihydro-

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Comment

1,2,4-Triazines and the compounds derived from them are found to possess a wide variety of pharmacological activities (Holla *et al.*, 1999). Triazine derivatives include herbicides having a broad spectrum of action that kill many species of weeds, and also herbicides with a narrow selectivity (Gruzdyev *et al.*, 1983). The title compound, (I), is an intermediate in the synthesis of a herbicide, 4-amino-6-(*tert*-butyl)-3-methylthio-1,2,4-triazin-5(4*H*)-one, commonly known as 'Sencor' (Eue & Tietz, 1972). It is used for the control of weeds in potato crops and is a broad-spectrum herbicide used in pre-emergence application in potato crops at a rate of 0.75–1 kg per hectare. In view of the importance of the title compound, (I), its crystal structure is reported here.



In all essential details, the molecular geometry (Fig. 1 and Table 1) is in good agreement with related structures (Ghassemzadeh *et al.*, 1998). The triazine ring is planar and exists in the thione form; the C=S bond length of 1.663 (2) Å is slightly longer than the pure double-bond distance (1.61 Å; Pauling, 1960). The N-N [1.341 (2) Å] and C-N [mean value 1.341 (2) Å] bond distances are intermediate between the expected single (1.45 and 1.47 Å, respectively) and double (1.20 and 1.27 Å, respectively) bond distances.

The crystal structure is stabilized by intra- and intermolecular hydrogen bonds (Table 2). Atom N1 of the triazine ring links the carbonyl O atom of an adjacent triazine ring, forming a chain running parallel to the c axis. The tertiary butyl group is oriented such that there are alternate hydrophobic and hydrophilic layers in the crystal packing (Fig. 2).

Experimental

The title compound, (I), was obtained as a gift sample from Rallies India Ltd, Bangalore. The compound was recrystallized by slow evaporation of an acetone–toluene (9:1) solution (m.p. 488 K).

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Figure 1

A view of (I), with the atom-numbering scheme. Displacement ellipsoids are drawn at the 50% probability level and H atoms are shown as small spheres of arbitrary radii.



Figure 2

A packing diagram of (I), viewed down the b axis, showing the chain formation. Dashed lines indicate N-H···O hydrogen bonds. H atoms not involved in hydrogen bonding have been omitted for clarity.

Crystal data

C7H12N4OS	$D_x = 1.308 \text{ Mg m}^{-3}$
$M_r = 200.27$	Mo $K\alpha$ radiation
Monoclinic, $P2_1/c$	Cell parameters from 6656
a = 12.5196 (8) Å	reflections
b = 6.9535 (4) Å	$\theta = 3.0-27.9^{\circ}$
c = 12.5987(8) Å	$\mu = 0.29 \text{ mm}^{-1}$
$\beta = 112.039 \ (1)^{\circ}$	T = 273 (2) K
$V = 1016.64 (11) \text{ Å}^3$	Block, colorless
Z = 4	$0.21 \times 0.17 \times 0.09 \text{ mm}$

Data collection

Bruker SMART APEX CCD area-	1640 reflections with $I > 2\sigma(I)$
detector diffractometer	$R_{\rm int} = 0.018$
ω scan	$\theta_{\rm max} = 25.0^{\circ}$
Absorption correction: none	$h = -14 \rightarrow 14$
9276 measured reflections	$k = -8 \rightarrow 8$
1780 independent reflections	$l = -14 \rightarrow 14$

Refinement

Refinement on F^2	$w = 1/[\sigma^2(F_0^2) + (0.0662P)^2]$
$R[F^2 > 2\sigma(F^2)] = 0.039$	+ 0.2933 <i>P</i>]
$vR(F^2) = 0.115$	where $P = (F_0^2 + 2F_c^2)/3$
S = 1.07	$(\Delta/\sigma)_{\rm max} < 0.001$
780 reflections	$\Delta \rho_{\rm max} = 0.40 \ {\rm e} \ {\rm \AA}^{-3}$
21 parameters	$\Delta \rho_{\rm min} = -0.35 \text{ e} \text{ \AA}^{-3}$
H-atom parameters constrained	

Table 1

Selected geometric parameters (Å, °).

C1-N1	1.336 (3)	C2-N2	1.378 (2)
C1-N2	1.370 (2)	C3-N3	1.288 (2)
2-01	1.2244 (19)	N2-N4	1.401 (2)

C1-N1-N3 127.70 (15)

Table 2		
Hydrogen-bond geometry	(Å.	°).

$D - H \cdot \cdot \cdot A$	D-H	$H \cdots A$	$D{\cdots}A$	$D - \mathbf{H} \cdots A$
N1-H1···O1 ⁱ	0.86	1.94	2.790 (2)	168
$N4 - H4B \cdot \cdot \cdot O1$	0.86	2.31	2.607 (2)	100
$N4-H4A\cdots S1$	0.86	2.59	2.976 (2)	108

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

All H atoms were included in calculated positions (C-H = 0.93-0.98 Å and N-H = 0.86 Å) and refined using a riding model, with $U_{\rm iso}({\rm H}) = 1.2 U_{\rm eq}({\rm N},C)$ or $1.5 U_{\rm eq}({\rm methyl C})$. The methyl groups were allowed to rotate but not to tip.

Data collection: SMART (Bruker, 2001); cell refinement: SAINT (Bruker, 2001); data reduction: SAINT; program(s) used to solve structure: SHELXS97 (Sheldrick, 1997); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: SHELXTL/PC (Sheldrick, 1990) and MERCURY (Bruno et al., 2002); software used to prepare material for publication: SHELXL97.

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