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Key indicators

Single-crystal X-ray study $T=293~{\rm K}$ Mean $\sigma({\rm C-C})=0.016~{\rm \AA}$ R factor = 0.097 wR factor = 0.231 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.

2-(1,1,2,2-Tetrachloroethylsulfanyl)-3a,4,7,7a-tetrahydro-1*H*-isoindole-1,3(2*H*)-dione

The title compound (also known as captafol), $C_{10}H_9Cl_4NO_2S$, exhibits fungicidal activity. The six-membered ring exists in a boat conformation. The crystal packing is stabilized by weak intermolecular $C-H\cdots O$ and $C-H\cdots Cl$ interactions.

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Comment

An important aspect of the rational design of bioactive molecules involves the search for structure–activity relationships (Lewis *et al.*, 1991). The conformation of the molecule is found to influence the level of biological activity of the proper compound (Baughman & Jacobson, 1978). The correlations found between the crystal structures of some compounds and their biological activity have aided in the chemical design of few active agrochemicals (Rohrbaugh & Jacobson, 1978). Earlier, the activity of a series of triazolyl ketone herbicides (Anderson *et al.*, 1983) has been investigated along with the fungicidal activities of *N*-phenylsuccinamides (Zenei *et al.*, 1988).

We present here the crystal structure of the title compound, (I) (Fig. 1), which posesses fungicidal activity. Atoms H4 and H5 at the fused ring junction have a *cis* relationship. The puckering parameters (Cremer & Pople, 1975), generated by *PLATON* (Spek, 2003), for the six-membered cyclohexene ring are $q_2 = 0.557$ (13) Å, $\varphi_2 = 117.3$ (12)° and $\tau = 91.7$ (12)°, indicating a boat conformation. The crystal structure is stabilized by weak intermolecular $C-H\cdots O$ and $C-H\cdots Cl$ interactions (Table 1) which link the molecules into linear chains along the *c* axis (Fig. 2).

Experimental

The title compound was obtained from Rallis India, Bangalore. Single crystals were grown by slow evaporation of a mixture of ethyl acetate/benzene at 278 K.

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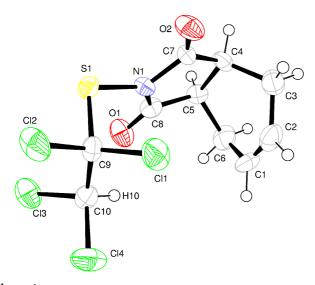


Figure 1 View of (I), shown with 50% probability displacement ellipsoids.

Crystal data

C ₁₀ H ₉ Cl ₄ NO ₂ S	$D_x = 1.683 \text{ Mg m}^{-3}$
$M_r = 349.04$	Mo $K\alpha$ radiation
Monoclinic, $P2_1/c$	Cell parameters from 675
a = 10.567 (8) Å	reflections
b = 6.674 (5) Å	$\theta = 2.2 – 26.4^{\circ}$
c = 19.427 (15) Å	$\mu = 1.01 \text{ mm}^{-1}$
$\beta = 91.715 (12)^{\circ}$	T = 293 (2) K
$V = 1369.6 (18) \text{ Å}^3$	Rod, colorless
Z = 4	$0.22 \times 0.07 \times 0.06 \text{ mm}$

Data collection

Bruker SMART APEX CCD area-	2449 independent reflections
detector diffractometer	2158 reflections with $I > 2\sigma(I)$
φ and ω scans	$R_{\rm int} = 0.039$
Absorption correction: multi-scan	$\theta_{\rm max} = 25.4^{\circ}$
(SADABS; Sheldrick, 1996)	$h = -11 \rightarrow 12$
$T_{\min} = 0.809, T_{\max} = 0.942$	$k = -8 \rightarrow 8$
8949 measured reflections	$l = -23 \rightarrow 23$

Refinement

Refinement on F^2	$w = 1/[\sigma^2(F_o^2) + (0.0809P)^2]$
$R[F^2 > 2\sigma(F^2)] = 0.097$	+ 27.5824 <i>P</i>]
$wR(F^2) = 0.231$	where $P = (F_o^2 + 2F_c^2)/3$
S = 1.17	$(\Delta/\sigma)_{\rm max} < 0.001$
2449 reflections	$\Delta \rho_{\text{max}} = 1.05 \text{ e Å}^{-3}$
167 parameters	$\Delta \rho_{\min} = -0.53 \text{ e Å}^{-3}$
H-atom parameters constrained	

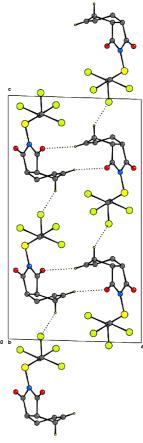
Table 1 Intermolecular contacts (Å, °).

$D-\mathrm{H}\cdots A$	D-H	$H \cdot \cdot \cdot A$	$D \cdot \cdot \cdot A$	$D-\mathrm{H}\cdots A$
$ \begin{array}{c} C2-H2\cdots O2^{i} \\ C3-H3A\cdots C12^{ii} \end{array} $	0.93	2.65	3.545 (15)	163 (1)
	0.97	2.88	3.616 (13)	134 (1)

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

All H atoms were placed in idealized positions (0.93–0.98 Å) and refined as riding, with $U_{\rm iso}({\rm H}) = 1.2 U_{\rm eq}({\rm C})$. The maximum positive peak in the difference Fourier map was located 1.15 Å from atom S1.

Data collection: *SMART* (Bruker, 2000); cell refinement: *SAINT* (Bruker, 2000); data reduction: *SAINT*; program(s) used to solve structure: *SIR*92 (Altomare *et al.*, 1993); program(s) used to refine



Packing diagram of (I), showing the intermolecular $C-H\cdots O$ and $C-H\cdots C$ 1 contacts (dashed lines).

structure: *SHELXL*97 (Sheldrick, 1997); molecular graphics: *ORTEP-3 for Windows* (Farrugia, 1997) and *CAMERON* (Watkin *et al.*, 1993); software used to prepare material for publication: *PLATON* (Spek, 2003).

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