Acta Crystallographica Section E Structure Reports Online

ISSN 1600-5368

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#### **Key indicators**

Single-crystal X-ray study T = 100 KMean  $\sigma$ (C–C) = 0.007 Å R factor = 0.070 wR factor = 0.223 Data-to-parameter ratio = 18.2

For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e.

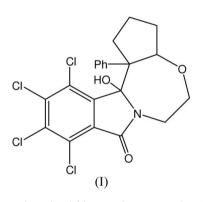
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# 9,10,11,12-Tetrachloro-12b-hydroxy-12c-phenyl-1,2,3,3a,5,6,12b,12c-octahydro-8*H*-cyclopenta[6,7][1,4]oxazepino[5,4-*a*]isoindol-8-one

The medium-sized heterocyclic title compound,  $C_{21}H_{17}Cl_4NO_3$ , crystallizes with two crystallographically independent molecules in the asymmetric unit. The cyclopentane rings adopt half-chair conformations. Symmetry-related molecules are linked *via* O-H···O hydrogen bonds to form dimers, which are arranged into one-dimensional chains along the [100] direction. C-H··· $\pi$  and  $\pi$ - $\pi$  interactions were also observed in the crystal structure.

### Comment

The construction of medium and large heterocyclic ring systems is an important task in organic synthesis (Evans & Holmes, 1991; Griesbeck *et al.*, 2001; Illuminati & Mandolini, 1981). In a continuation of our recent work on photoinduced cyclization reactions (Xue *et al.*, 2000), we prepared the title compound, (I), by the photochemical reaction of N-(2-hydroxyethyl)-4,5,6,7-tetrachlorophthalimide with 1-phenyl-cyclopentane and we report its structure here.



The asymmetric unit of (I) contains two molecules, A and B, that differ little in their conformations (Fig. 1). Their bond lengths and angles are similar and show normal values (Allen *et al.*, 1987) that are comparable to related structures (Xue *et al.*, 2000). In both molecules, the cyclopentane rings adopt half-chair conformations, with puckering parameters (Cremer & Pople 1975) Q = 0.426 (6) Å and  $\varphi = 95.9$  (7)° in molecule A [Q = 0.449 (5) Å and  $\varphi = 141.8$  (7)° in molecule B]. The conformation of the seven-membered ring is twist-chair. The dihedral angles between the C16–C21 benzene rings and the mean planes of the C1–C8/N1 units are 31.9 (2)° in molecule A and 39.6 (2)° in molecule B.

The crystal packing of (I) (Fig. 2) shows that adjacent molecules are linked into dimers by  $O3A-H3AA\cdots O2B$  hydrogen bonds (Table 1) and these dimers are arranged into one-dimensional chains along the [100] direction. The molecular structure is further stabilized by  $C-H\cdots\pi$  interactions (Table 1).  $\pi-\pi$  Interactions are also observed with centroid–

Received 18 December 2006 Accepted 21 December 2006 centroid distances of  $Cg1\cdots Cg5 = 3.444$  (3) Å,  $Cg3\cdots Cg5 = 3.594$  (3) Å,  $Cg2\cdots Cg6 = 3.839$  (3) Å and  $Cg4\cdots Cg6 = 3.725$  (3) Å, where Cg1 is the centroid of the C1A/C2A/C7A/C8A/N1A ring, Cg2 that of C1B/C2B/C7B/C8B/N1B; Cg3, Cg4, Cg5 and Cg6 those of the C2A-C7A, C2B-C7B, C16A-C21A and C16B-C21B benzene rings, respectively.

## Experimental

Compound (I) was synthesized by a photoinduced reaction between N-(2-hydroxyethyl)-4,5,6,7-tetrachlorophthalimide (3 mmol) and an excess of 1-phenylcyclopentene (15 mmol) in a benzene solution (120 ml), using a medium-pressure mercury lamp in a glass cooling water jacket as the light source. It was isolated by column chromatography on silica gel using petroleum ether and ethyl acetate as eluants. Colorless block-shaped single crystals suitable for X-ray diffraction analysis were obtained by slow evaporation of the solvents from a petroleum ether/chloroform solution (1:1 v/v), m.p. 538–540 K.

 $V = 2006.26 (14) \text{ Å}^3$ 

 $D_x = 1.566 \text{ Mg m}^{-3}$ Mo  $K\alpha$  radiation  $\mu = 0.61 \text{ mm}^{-1}$ T = 100.0 (1) KBlock, colorless  $0.59 \times 0.47 \times 0.24 \text{ mm}$ 

29592 measured reflections

 $R_{\rm int} = 0.031$ 

 $\theta_{\rm max} = 28.0^\circ$ 

9537 independent reflections 8288 reflections with  $I > 2\sigma(I)$ 

Z = 4

### Crystal data

C <sub>21</sub> H <sub>17</sub> Cl <sub>4</sub> NO <sub>3</sub>
$M_r = 473.16$
Triclinic, P1
a = 9.0183 (4)  Å
b = 12.5563 (5)  Å
c = 18.3224 (7) Å
$\alpha = 75.391 \ (2)^{\circ}$
$\beta = 88.559 \ (2)^{\circ}$
$\gamma = 88.083 \ (2)^{\circ}$

Data collection

Bruker SMART APEX2 CCD areadetector diffractometer  $\omega$  scans Absorption correction: multi-scan (*SADABS*; Bruker, 2005)  $T_{\rm min} = 0.712, T_{\rm max} = 0.867$ 

### Refinement

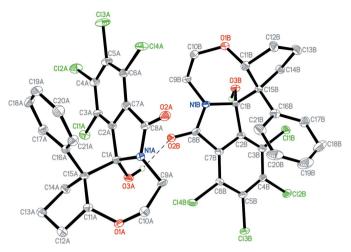
Refinement on $F^2$	$w = 1/[\sigma^2(F_o^2) + (0.0783P)^2]$
$R[F^2 > 2\sigma(F^2)] = 0.070$	+ 15.8392P]
$wR(F^2) = 0.223$	where $P = (F_0^2 + 2F_c^2)/3$
S = 1.15	$(\Delta/\sigma)_{\rm max} < 0.001$
9537 reflections	$\Delta \rho_{\rm max} = 1.44 \text{ e } \text{\AA}^{-3}$
523 parameters	$\Delta \rho_{\rm min} = -0.68 \text{ e } \text{\AA}^{-3}$
H-atom parameters constrained	

Table	1
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Hydrogen-bond geometry (Å, °).

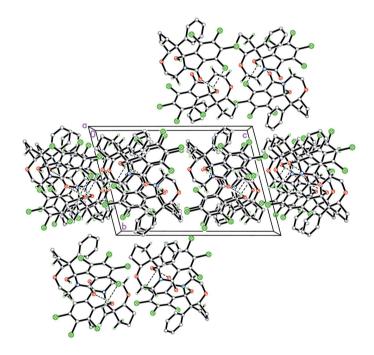
$D - H \cdot \cdot \cdot A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdot \cdot \cdot A$
$O3A - H3AA \cdots O2B$	0.82	1.97	2.763 (5)	162
$O3B-H3BA\cdots O2A^{i}$	0.82	1.94	2.751 (5)	173
$C9A - H9AA \cdots O2A$	0.97	2.45	2.852 (5)	104
$C9B - H9BA \cdots O2B$	0.97	2.48	2.870 (5)	104
C9 <i>B</i> −H9 <i>BB</i> ···O3 <i>B</i>	0.97	2.52	2.875 (6)	102
$C9B - H9BB \cdots O1B^{ii}$	0.97	2.59	3.357 (5)	136
$C10A - H10A \cdots O1A^{iii}$	0.97	2.59	3.096 (6)	113
$C14A - H14A \cdots Cl1A$	0.97	2.68	3.292 (5)	121
$C14A - H14B \cdots O3A$	0.97	2.37	2.796 (6)	106
$C14B - H14D \cdots O3B$	0.97	2.32	2.675 (6)	101
$C9A - H9AA \cdots Cg4^{iv}$	0.97	3.25	3.938 (5)	130

Symmetry codes: (i) x - 1, y, z; (ii) -x, -y + 1, -z + 1; (iii) -x + 1, -y + 1, -z; (iv) x + 1, y, z.





The asymmetric unit of (I), showing 50% probability displacement ellipsoids and the atomic numbering. For clarity, only hydroxyl H atoms are shown, with the hydrogen bond drawn as a dashed line.



### Figure 2

The crystal packing of (I), viewed down the a axis. Hydrogen bonds are drawn as dashed lines. H atoms not involved in these interactions have been omitted.

All H-atoms were positioned geometrically and refined using a riding model with C-H = 0.93 Å for aromatic, 0.98 Å for CH and 0.97 Å for CH<sub>2</sub>, and O-H = 0.82 Å;  $U_{\rm iso}({\rm H}) = 1.2U_{\rm eq}({\rm C})$  or  $1.5U_{\rm eq}({\rm O})$ . The highest residual electron density peak is located 0.92 Å from Cl1A.

Data collection: *APEX2* (Bruker, 2005); cell refinement: *APEX2*; data reduction: *SAINT* (Bruker, 2005); program(s) used to solve structure: *SHELXTL* (Sheldrick, 1998); program(s) used to refine structure: *SHELXTL*; molecular graphics: *SHELXTL*; software used to prepare material for publication: *SHELXTL* and *PLATON* (Spek, 2003).

This work was supported by the National Natural Science Foundation of China and the Natural Science Foundation of Jiangsu Province. The authors also thank the Malaysian Government and Universiti Sains Malaysia for the Scientific Advancement Grant Allocation (SAGA) grant No. 304/ PFIZIK/653003/A118.

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