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Key indicators

Single-crystal X-ray study T = 293 K Mean σ (C–C) = 0.003 Å R factor = 0.036 wR factor = 0.104 Data-to-parameter ratio = 15.4

For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e. The asymmetric unit of the title compound, $C_8H_3ClO_3$, contains two molecules. The molecular packing is controlled by $C-H \cdots O$ hydrogen bonding.

5-Chloro-2-benzofuran-1,3-dione

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Comment

The title compound, (I), is a valuable intermediate for Cu phthalocyanine and anthraquinone dyes (Bansho *et al.*, 1960), quinophthalone pigments (Dietmar *et al.*, 1998) and polyether–polyimide polymers (Brunelle *et al.*, 1999). Compound (I) can be made from phthalic anhydride by chlorination, but it is very difficult to separate (I) from the by-product 4,5dichlorophthalic anhydride by recrystallization. This paper reports the structure analysis of (I).



The asymmetric unit contains two molecules, which are very similar (Fig. 1). These two molecules are essentially planar, making a dihedral angle of 58.13 (5)°. Except for the C–Cl bond lengths, which are 0.01 Å longer than those observed in 4,5-dichlorophthalic anhydride (Ojala *et al.*, 1999), other bond lengths and angles are similar for the two compounds. The packing of the molecules in the crystal is governed by weak C–H···O hydrogen bonds (Fig. 2 and Table 2).

Experimental

The title compound was obtained by dehydration of 4-chlorophthalic acid, which was synthesized in our laboratory following the procedure described by Ayling (1929). Crystals suitable for X-ray analysis were obtained by slow evaporation of a saturated carbon tetrachloride solution at room temperature.



Figure 1

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The asymmetric unit of (I), showing the atomic numbering. Displacement ellipsoids are drawn at the 50% probability level for non-H atoms.



Figure 2

Packing diagram of (I), showing the hydrogen bonds as dashed lines.

Crystal data

C ₈ H ₃ ClO ₃	Z = 4
$M_r = 182.55$	$D_x = 1.604 \text{ Mg m}^{-3}$
Triclinic, $P\overline{1}$	Mo $K\alpha$ radiation
a = 6.929 (2) Å	Cell parameters from 4173
$b = 7.8540 (10) \text{\AA}$	reflections
c = 14.262 (2) Å	$\theta = 1.5-27.2^{\circ}$
$\alpha = 77.69 \ (1)^{\circ}$	$\mu = 0.46 \text{ mm}^{-1}$
$\beta = 88.09 \ (1)^{\circ}$	T = 293 (2) K
$\gamma = 85.48 \ (2)^{\circ}$	Prism, colourless
$V = 755.8 (3) \text{ Å}^3$	$0.4 \times 0.2 \times 0.1 \text{ mm}$

 $\begin{array}{l} R_{\mathrm{int}}=0.015\\ \theta_{\mathrm{max}}=27.2^\circ\\ h=-1\rightarrow8\\ k=-10\rightarrow10\\ l=-18\rightarrow18 \end{array}$

3 standard reflections

every 100 reflections

intensity decay: 0.5%

 $w = 1/[\sigma^2(F_o^2) + (0.0367P)^2]$

where $P = (F_o^2 + 2F_c^2)/3$

+ 0.2624P]

 $(\Delta/\sigma)_{\rm max} = 0.001$

 $\Delta \rho_{\rm max} = 0.25 \text{ e} \text{ \AA}^{-3}$

 $\Delta \rho_{\rm min} = -0.24 \text{ e} \text{ Å}^{-3}$

Data collection

Enraf-Nonius CAD-4
diffractometer
$\omega/2\theta$ scans
Absorption correction: ψ scan
(North et al., 1968)
$T_{\min} = 0.837, T_{\max} = 0.955$
4173 measured reflections
3336 independent reflections
2184 reflections with $I > 2\sigma(I)$

Refinement

Refinement on F^2 $R[F^2 > 2\sigma(F^2)] = 0.036$ $wR(F^2) = 0.104$ S = 1.083336 reflections 217 parameters H-atom parameters constrained

Table 1

Selected geometric parameters (Å).

C1-C2	1.380 (3)	C9-C10	1.374 (3)
C1-C7	1.468 (3)	C9-C15	1.470 (3)
C2-C8	1.470 (3)	C10-C16	1.471 (3)
C4-Cl2	1.729 (2)	C12-Cl1	1.731 (2)
C7-O1	1.189 (3)	C15-O4	1.193 (3)
C7-O2	1.388 (3)	C15-O5	1.386 (3)
C8-O3	1.183 (3)	C16-O6	1.181 (3)
C8-O2	1.393 (3)	C16-O5	1.395 (3)

Table 2		
Hydrogen-bonding geometry	(Å,	°).

$D - H \cdot \cdot \cdot A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdot \cdot \cdot A$
C3-H3···O1 ⁱ	0.93	2.80	3.292 (3)	114
C6-H6···O6 ⁱⁱ	0.93	3.03	3.466 (3)	111
C6−H6···O6 ⁱⁱⁱ	0.93	2.89	3.690 (3)	145
$C11 - H11 \cdots O2^{iv}$	0.93	2.94	3.675 (3)	137
C13-H13···O1	0.93	2.63	3.522 (3)	161
C13-H13···O2	0.93	2.81	3.453 (3)	127
$C14-H14\cdots O4^{v}$	0.93	2.61	3.519 (3)	166

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

All H atoms were located in difference Fourier maps but were introduced in the refinement in calculated positions (C-H = 0.93 Å) and treated as riding, with $U_{iso}(H) = 1.2U_{eq}(C)$

Data collection: *CAD-4 EXPRESS* (Enraf–Nonius, 1994; cell refinement: *CAD-4 EXPRESS*; data reduction: *XCAD4* (Harms & Wocadlo, 1995); program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *ORTEP-3 for Windows* (Farrugia, 1997); software used to prepare material for publication: *WinGX* (Farrugia, 1999).

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