

2-Bromobenz[*a*]anthracene-5,6,7,12-tetraone-5-dimethylketal

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## Key indicators

Single-crystal X-ray study

$T = 293\text{ K}$

Mean  $\sigma(\text{C}-\text{C}) = 0.006\text{ \AA}$

$R$  factor = 0.066

$wR$  factor = 0.165

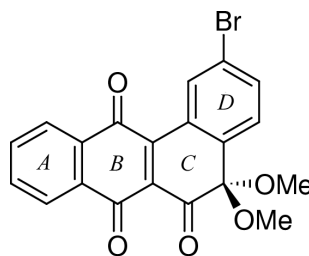
Data-to-parameter ratio = 16.5

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

In the title compound, 2-bromo-5,5-dimethoxybenzo[*a*]anthracene-6,7,12(5*H*)trione,  $\text{C}_{20}\text{H}_{13}\text{BrO}_5$ , the two aromatic rings are essentially planar and make a dihedral angle of  $15.1(2)^\circ$  with one another, while the dimethoxy-substituted cyclohexane ring is non-planar, as expected. An intramolecular  $\text{C}-\text{H}\cdots\text{O}$  interaction is present, with  $\text{C}\cdots\text{O} = 2.814(5)\text{ \AA}$  and  $\text{C}-\text{H}\cdots\text{O} = 125^\circ$ .

## Comment

The analogues of angucyclines have been intensively investigated due to their biological properties (Krohn & Rohr, 1997). In order to generate a new analogue, we have prepared the title compound, (I). We have undertaken an X-ray structural determination of compound (I) to define the conformation of this system, especially the conformation of the bromophenyl with respect to the substituted anthracene moiety.



(I)

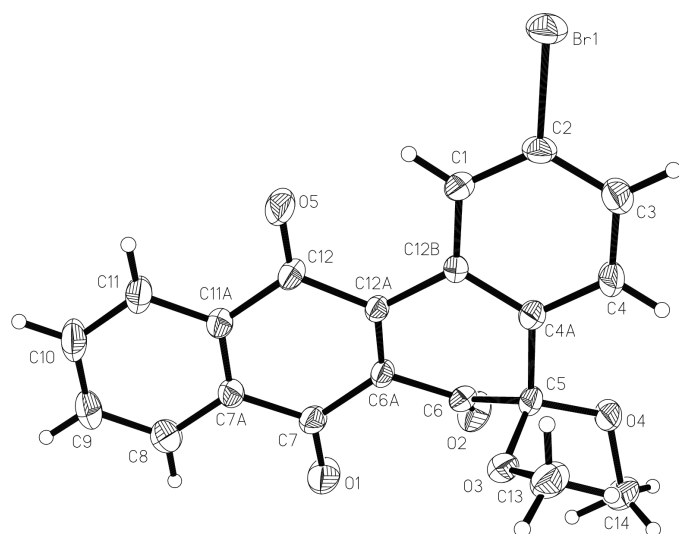
In (I), the bond lengths and angles have normal values. In the aromatic rings *A* and *D*, the average  $\text{C}-\text{C}$  bond distances are  $1.387$  and  $1.395\text{ \AA}$ , respectively. In comparison, the  $\text{C}-\text{C}$  bond distances within the cyclohexanedione (ring *B*) and that of  $\text{C}6\text{A}-\text{C}12\text{A}$  are slightly shorter than in the related structure, benz[*a*]anthracene-7,12-dione, (II), previously studied (Kuroda *et al.*, 1982), while the observed intramolecular interaction between  $\text{C}1\text{A}-\text{H}1\text{A}$  and  $\text{O}5$  in (I) is slightly longer than in (II) ( $2.058\text{ \AA}$ ).

The aromatic rings *A* and *D* are essentially planar, and atom  $\text{Br}1$  deviates slightly (by  $0.035\text{ \AA}$ ) from the least-squares plane through ring *D*. The angle between these two aromatic rings is  $15.1(2)^\circ$  and is much larger than in (II) benz[*a*]anthracene-7,12-dione (Kuroda *et al.*, 1982), because in (I), ring *C*, bridging between rings *A* and *D*, is non-planar. The close interaction between  $\text{O}5$  and  $\text{H}1\text{A}$ , and the non-bonded lone-pair-lone-pair repulsions of the  $\text{O}$  atoms [the non-bonded distance between  $\text{O}1$  and  $\text{O}2$  is  $2.880(5)\text{ \AA}$ ] are two reasons for the distortions from planarity. Atom  $\text{O}1$  lies on, while atom  $\text{O}5$

Received 23 July 2001

Accepted 31 July 2001

Online 10 August 2001



**Figure 1**  
The structure of the title compound showing 50% probability displacement ellipsoids and the atom-numbering scheme.

deviates slightly by  $-0.121(4)$  Å from the least-squares plane through ring *B* having a maximum deviation of  $0.052(4)$  Å (C12A). Atom O2 attached to ring *C* at C6 significantly deviates by  $-1.087(4)$  Å, while atom C6 deviates by  $-0.317(4)$  Å from the least-squares plane through ring *C*. The puckering parameters of ring *C* (Cremer & Pople, 1975), which adopts a half-chair conformation, are  $Q_T = 0.478(5)$  Å,  $\theta = 115.5(6)^\circ$  and  $\varphi_2 = 275.5(6)^\circ$ .

## Experimental

To a solution of 2-bromo-6-hydroxy-5-methoxybenz[a]anthracene in methanol was added an equimolar amount of phenyliodonium acetate. The mixture was stirred for 3 h at room temperature. After stirring, the solvent was removed *in vacuo*, and the residue was purified by column chromatography. The residue afforded a brown-red solid and was recrystallized from a mixture of ethyl acetate and petroleum ether (25/75).

### Crystal data

$C_{20}H_{13}BrO_5$   
 $M_r = 413.21$   
Triclinic,  $P\bar{1}$   
 $a = 7.7414(1)$  Å  
 $b = 10.3084(1)$  Å  
 $c = 10.9583(2)$  Å  
 $\alpha = 74.415(1)^\circ$   
 $\beta = 78.835(1)^\circ$   
 $\gamma = 81.192(1)^\circ$   
 $V = 821.65(2)$  Å<sup>3</sup>

### Data collection

Siemens SMART CCD diffractometer  
 $\omega$  scans  
Absorption correction: empirical (SADABS; Sheldrick, 1996)  
 $T_{\min} = 0.463$ ,  $T_{\max} = 0.786$   
5992 measured reflections

$Z = 2$   
 $D_x = 1.670$  Mg m<sup>-3</sup>  
Mo  $K\alpha$  radiation  
Cell parameters from 3602 reflections  
 $\theta = 2.0$ – $28.3^\circ$   
 $\mu = 2.53$  mm<sup>-1</sup>  
 $T = 293(2)$  K  
Slab, brown-red  
 $0.36 \times 0.26 \times 0.10$  mm

3910 independent reflections  
2584 reflections with  $I > 2\sigma(I)$   
 $R_{\text{int}} = 0.055$   
 $\theta_{\text{max}} = 28.3^\circ$   
 $h = -10 \rightarrow 10$   
 $k = -12 \rightarrow 13$   
 $l = -14 \rightarrow 10$

### Refinement

Refinement on  $F^2$   
 $R[F^2 > 2\sigma(F^2)] = 0.066$   
 $wR(F^2) = 0.165$   
 $S = 0.94$   
3910 reflections  
237 parameters

H-atom parameters constrained  
 $w = 1/[\sigma^2(F_o^2) + (0.0817P)^2]$   
where  $P = (F_o^2 + 2F_c^2)/3$   
 $(\Delta/\sigma)_{\text{max}} < 0.001$   
 $\Delta\rho_{\text{max}} = 0.97$  e Å<sup>-3</sup>  
 $\Delta\rho_{\text{min}} = -1.79$  e Å<sup>-3</sup>

**Table 1**

Selected geometric parameters (Å, °).

Br1–C2	1.896 (4)	C4–C4A	1.402 (6)
O1–C7	1.216 (5)	C1–C12B	1.400 (6)
O2–C6	1.200 (5)	C4A–C12B	1.416 (5)
O3–C5	1.430 (5)	C4A–C5	1.525 (5)
O4–C5	1.388 (5)	C5–C6	1.534 (6)
O5–C12	1.208 (5)	C6–C6A	1.499 (5)
C1–C2	1.386 (6)	C6A–C12A	1.353 (6)
C2–C3	1.385 (6)	C7A–C11A	1.388 (6)
C3–C4	1.379 (6)	C12A–C12B	1.498 (5)
C5–O3–C13	116.3 (3)	O3–C5–O4	112.9 (3)
C5–O4–C14	117.3 (3)	O2–C6–C5	123.4 (4)
C1–C2–C3	121.7 (4)	O2–C6–C6A	124.1 (4)
C1–C2–Br1	120.4 (3)	C5–C6–C6A	112.5 (3)
C3–C2–Br1	118.0 (3)	C12B–C4A–C5–C6	33.5 (5)
C12B–C4A–C5–C6	33.5 (5)	C11–C11A–C12–O5	−4.4 (7)
O4–C5–C6–O2	3.7 (6)	C11A–C12–C12A–C12B	−173.0 (4)
O2–C6–C6A–C7	47.0 (6)	C12–C12A–C12B–C1	−13.3 (6)

**Table 2**

Hydrogen-bonding geometry (Å, °).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
C1–H1A $\cdots$ O5	0.93	2.18	2.814 (5)	125

After checking their presence in the difference map, all H atoms were geometrically fixed and allowed to ride on their attached atoms.

Data collection: SMART (Siemens, 1996); cell refinement: SAINT (Siemens, 1996); data reduction: SAINT; program(s) used to solve structure: SHELXTL (Sheldrick, 1997); program(s) used to refine structure: SHELXTL; molecular graphics: SHELXTL; software used to prepare material for publication: SHELXTL, PARST (Nardelli, 1995) and PLATON (Spek, 1990).

The authors would like to thank the Malaysian Government and Universiti Sains Malaysia for research grant R&D No. 305/PFIZIK/610961, and also wish to thank CSIR, New Delhi, for financial support. SD would like to thank UGC for a fellowship and AU wishes to thank Universiti Sains Malaysia for a Visiting Postdoctoral Fellowship.

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