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(1*R*,1'*S*)-1,1'-Dihydroxy-1,1'-biisobenzofuran-3,3'(1*H*,1'*H*)-dione

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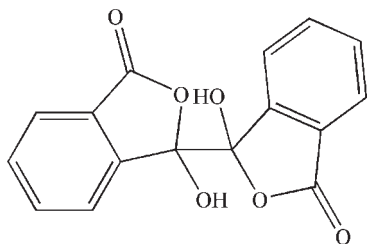
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 Key indicators: single-crystal X-ray study; $T = 293$ K; mean $\sigma(\text{C}-\text{C}) = 0.006$ Å; R factor = 0.056; wR factor = 0.170; data-to-parameter ratio = 10.4.

In the title compound, $\text{C}_{16}\text{H}_{10}\text{O}_6$, the complete molecule is generated by a crystallographic centre of symmetry. In the crystal, $\text{O}-\text{H}\cdots\text{O}$ hydrogen bonds link the molecules into (100) sheets and $\text{C}-\text{H}\cdots\text{O}$ links also occur.

Related literature

For background to phthalides as natural products, see: Pedrosa *et al.* (2006). For a related structure, see: Wang *et al.* (2001).



Experimental

Crystal data

 $\text{C}_{16}\text{H}_{10}\text{O}_6$
 $M_r = 298.24$

Monoclinic, $P2_1/c$
 $a = 8.2260$ (16) Å
 $b = 7.9690$ (16) Å
 $c = 10.859$ (4) Å
 $\beta = 114.03$ (2)°
 $V = 650.1$ (3) Å³

$Z = 2$
 Mo $K\alpha$ radiation
 $\mu = 0.12$ mm⁻¹
 $T = 293$ K
 $0.16 \times 0.12 \times 0.10$ mm

Data collection

Bruker SMART CCD diffractometer
 Absorption correction: none
 1352 measured reflections

1263 independent reflections
 622 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.070$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.056$
 $wR(F^2) = 0.170$
 $S = 1.02$
 1263 reflections
 121 parameters

H atoms treated by a mixture of independent and constrained refinement

$\Delta\rho_{\text{max}} = 0.29$ e Å⁻³
 $\Delta\rho_{\text{min}} = -0.30$ e Å⁻³

Table 1

Hydrogen-bond geometry (Å, °).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
$\text{O2}-\text{H2B}\cdots\text{O1}^{\text{i}}$	0.91 (7)	1.82 (7)	2.691 (5)	159 (5)
$\text{C5}-\text{H5A}\cdots\text{O1}^{\text{ii}}$	0.96 (3)	2.58 (4)	3.475 (5)	155 (3)

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

Data collection: *SMART* (Bruker, 1997); cell refinement: *SAINT* (Bruker, 1997); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *SHELXTL* (Sheldrick, 2008); software used to prepare material for publication: *SHELXTL*.

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: HB5170).

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supporting information

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(1*R*,1'*S*)-1,1'-Dihydroxy-1,1'-biisobenzofuran-3,3'(1*H*,1'*H*)-dione**Fang-Fang Jian, Shan-Shan Zhao, Huan-Mei Guo, Yu-Feng Li and Pu-Su Zhao****S1. Comment**

Substituted phthalides (isobenzofuran-1(3*H*)-ones) represent an important class of natural products that possess significant biological properties (e.g. Pedrosa *et al.*, 2006). As part of our search for new biologically active compounds, we unexpectedly obtained the title compound, (I), which is a typical derivative of phthalides.

In the crystal structure of compound (I) (Fig. 1), there is an inversion center, which is located at the mid-point of C(8)—C(8A) bond. All of the bond lengths and bond angles are in the normal ranges (Wang *et al.*, 2001). In the crystal lattice, there are a C—H···O intramolecular hydrogen bond and an O—H···O intermolecular hydrogen bond, which stabilize the molecule structure.

S2. Experimental

Phthalic anhydride (0.05 mol) was dissolved in dichloromethane (100 ml). Then, AlCl₃ (0.05 mol) was added. The mixture was stirred at room temperature and the whole reaction was under the protection of nitrogen. After 5 h, the reaction was stopped and the mixture poured into ice-water. The organic layer was collected and then was dried with MgSO₄. Finally, the organic layer was concentrated by rotary vacuum evaporation to obtain yellow solids. Yellow blocks of (I) were obtained by recrystallization from acetonitrile at room temperature.

S3. Refinement

The H atoms were located in difference maps and freely refined.

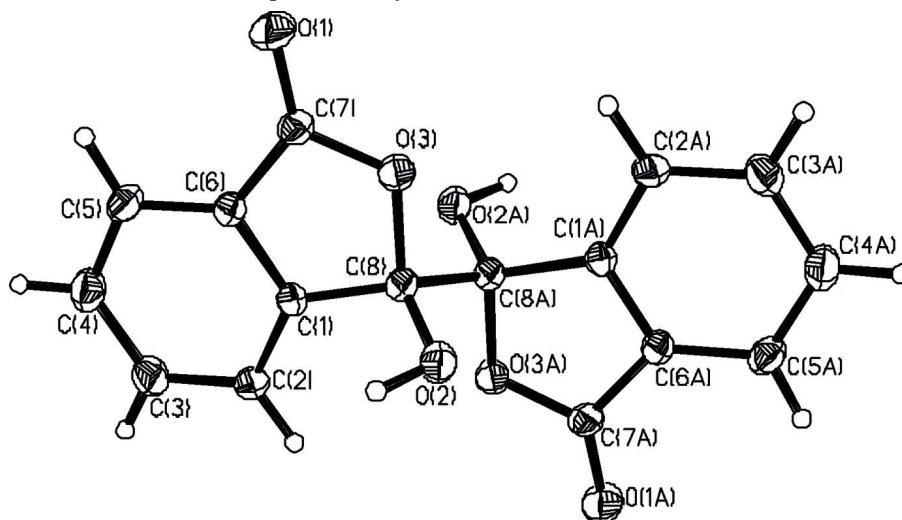


Figure 1

The molecular structure of (I) with displacement ellipsoids drawn at the 30% probability level.

(1*R*,1'*S*)-1,1'-dihydroxy-1,1'-biisobenzofuran- 3,3'(1*H*,1'*H*)-dione

Crystal data

$C_{16}H_{10}O_6$	$F(000) = 308$
$M_r = 298.24$	$D_x = 1.523 \text{ Mg m}^{-3}$
Monoclinic, $P2_1/c$	Mo $K\alpha$ radiation, $\lambda = 0.71073 \text{ \AA}$
Hall symbol: -P 2ybc	Cell parameters from 1978 reflections
$a = 8.2260 (16) \text{ \AA}$	$\theta = 3.5\text{--}27.5^\circ$
$b = 7.9690 (16) \text{ \AA}$	$\mu = 0.12 \text{ mm}^{-1}$
$c = 10.859 (4) \text{ \AA}$	$T = 293 \text{ K}$
$\beta = 114.03 (2)^\circ$	Block, yellow
$V = 650.1 (3) \text{ \AA}^3$	$0.16 \times 0.12 \times 0.10 \text{ mm}$
$Z = 2$	

Data collection

Bruker SMART CCD diffractometer	622 reflections with $I > 2\sigma(I)$
Radiation source: fine-focus sealed tube	$R_{\text{int}} = 0.070$
Graphite monochromator	$\theta_{\text{max}} = 25.9^\circ$, $\theta_{\text{min}} = 2.7^\circ$
ω scans	$h = 0 \rightarrow 9$
1352 measured reflections	$k = -9 \rightarrow 0$
1263 independent reflections	$l = -13 \rightarrow 12$

Refinement

Refinement on F^2	Hydrogen site location: inferred from neighbouring sites
Least-squares matrix: full	H atoms treated by a mixture of independent and constrained refinement
$R[F^2 > 2\sigma(F^2)] = 0.056$	$w = 1/[\sigma^2(F_o^2) + (0.0846P)^2]$
$wR(F^2) = 0.170$	where $P = (F_o^2 + 2F_c^2)/3$
$S = 1.02$	$(\Delta/\sigma)_{\text{max}} < 0.001$
1263 reflections	$\Delta\rho_{\text{max}} = 0.29 \text{ e \AA}^{-3}$
121 parameters	$\Delta\rho_{\text{min}} = -0.30 \text{ e \AA}^{-3}$
0 restraints	Extinction correction: <i>SHELXL97</i> (Sheldrick, 2008), $F_c^* = kFc[1 + 0.001x \text{Fc}^2\lambda^3/\sin(2\theta)]^{-1/4}$
Primary atom site location: structure-invariant direct methods	Extinction coefficient: 0.032 (10)
Secondary atom site location: difference Fourier map	

Special details

Geometry. All e.s.d.'s (except the e.s.d. in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell e.s.d.'s are taken into account individually in the estimation of e.s.d.'s in distances, angles and torsion angles; correlations between e.s.d.'s in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell e.s.d.'s is used for estimating e.s.d.'s involving l.s. planes.

Refinement. Refinement of F^2 against ALL reflections. The weighted R -factor wR and goodness of fit S are based on F^2 , conventional R -factors R are based on F , with F set to zero for negative F^2 . The threshold expression of $F^2 > \sigma(F^2)$ is used only for calculating R -factors(gt) *etc.* and is not relevant to the choice of reflections for refinement. R -factors based on F^2 are statistically about twice as large as those based on F , and R -factors based on ALL data will be even larger.

Fractional atomic coordinates and isotropic or equivalent isotropic displacement parameters (\AA^2)

	x	y	z	$U_{\text{iso}}^*/U_{\text{eq}}$
O1	0.2464 (4)	0.3665 (4)	0.1300 (3)	0.0549 (9)

O2	0.6822 (3)	0.5869 (4)	0.4778 (3)	0.0410 (8)
O3	0.4600 (3)	0.4127 (3)	0.3342 (2)	0.0376 (8)
C1	0.3733 (5)	0.6782 (5)	0.3741 (3)	0.0314 (9)
C2	0.3529 (6)	0.8340 (5)	0.4222 (4)	0.0398 (11)
C3	0.2204 (6)	0.9369 (6)	0.3366 (4)	0.0467 (11)
C4	0.1095 (6)	0.8864 (6)	0.2078 (4)	0.0484 (12)
C5	0.1271 (5)	0.7290 (6)	0.1605 (4)	0.0408 (11)
C6	0.2612 (5)	0.6276 (5)	0.2465 (3)	0.0316 (9)
C7	0.3138 (5)	0.4584 (5)	0.2258 (3)	0.0359 (10)
C8	0.5080 (5)	0.5441 (5)	0.4390 (3)	0.0330 (10)
H2A	0.435 (5)	0.868 (4)	0.511 (4)	0.036 (10)*
H4A	0.014 (5)	0.956 (6)	0.148 (4)	0.052 (12)*
H5A	0.052 (5)	0.686 (5)	0.073 (3)	0.036 (10)*
H3A	0.206 (6)	1.043 (7)	0.368 (4)	0.069 (15)*
H2B	0.689 (9)	0.669 (8)	0.422 (6)	0.12 (2)*

Atomic displacement parameters (Å²)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
O1	0.067 (2)	0.049 (2)	0.0364 (15)	0.0066 (16)	0.0091 (15)	-0.0153 (15)
O2	0.0378 (16)	0.0411 (18)	0.0391 (15)	-0.0008 (14)	0.0105 (13)	0.0080 (13)
O3	0.0498 (17)	0.0327 (16)	0.0273 (13)	0.0054 (13)	0.0125 (13)	-0.0038 (12)
C1	0.036 (2)	0.028 (2)	0.0265 (18)	-0.0023 (16)	0.0096 (17)	0.0042 (16)
C2	0.052 (3)	0.032 (2)	0.0280 (19)	-0.003 (2)	0.008 (2)	-0.0055 (18)
C3	0.057 (3)	0.035 (3)	0.046 (2)	0.010 (2)	0.018 (2)	0.000 (2)
C4	0.045 (3)	0.049 (3)	0.043 (2)	0.011 (2)	0.009 (2)	0.008 (2)
C5	0.039 (2)	0.048 (3)	0.028 (2)	0.000 (2)	0.0058 (18)	-0.003 (2)
C6	0.035 (2)	0.032 (2)	0.0263 (18)	0.0009 (17)	0.0107 (17)	0.0015 (16)
C7	0.044 (2)	0.037 (3)	0.0251 (19)	-0.0039 (19)	0.0119 (18)	-0.0032 (17)
C8	0.037 (2)	0.030 (2)	0.0262 (18)	0.0016 (18)	0.0074 (17)	-0.0005 (16)

Geometric parameters (Å, °)

O1—C7	1.207 (4)	C2—H2A	0.96 (4)
O2—C8	1.362 (4)	C3—C4	1.382 (6)
O2—H2B	0.91 (7)	C3—H3A	0.94 (5)
O3—C7	1.346 (4)	C4—C5	1.385 (6)
O3—C8	1.477 (4)	C4—H4A	0.96 (4)
C1—C6	1.376 (5)	C5—C6	1.380 (5)
C1—C2	1.383 (5)	C5—H5A	0.96 (4)
C1—C8	1.494 (5)	C6—C7	1.461 (5)
C2—C3	1.378 (6)	C8—C8 ⁱ	1.551 (7)
C8—O2—H2B	108 (4)	C6—C5—H5A	118 (2)
C7—O3—C8	110.2 (3)	C4—C5—H5A	125 (2)
C6—C1—C2	120.6 (4)	C1—C6—C5	122.2 (4)
C6—C1—C8	109.2 (3)	C1—C6—C7	107.9 (3)
C2—C1—C8	130.2 (3)	C5—C6—C7	129.9 (3)

C3—C2—C1	117.6 (4)	O1—C7—O3	121.5 (4)
C3—C2—H2A	123 (2)	O1—C7—C6	129.2 (4)
C1—C2—H2A	119 (2)	O3—C7—C6	109.3 (3)
C2—C3—C4	121.6 (5)	O2—C8—O3	109.5 (3)
C2—C3—H3A	118 (3)	O2—C8—C1	116.8 (3)
C4—C3—H3A	120 (3)	O3—C8—C1	103.2 (3)
C3—C4—C5	120.9 (4)	O2—C8—C8 ⁱ	107.1 (4)
C3—C4—H4A	122 (2)	O3—C8—C8 ⁱ	104.4 (4)
C5—C4—H4A	117 (2)	C1—C8—C8 ⁱ	115.0 (4)
C6—C5—C4	117.1 (4)		

Symmetry code: (i) $-x+1, -y+1, -z+1$.

Hydrogen-bond geometry (\AA , $^\circ$)

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
O2—H2B \cdots O1 ⁱⁱ	0.91 (7)	1.82 (7)	2.691 (5)	159 (5)
C5—H5A \cdots O1 ⁱⁱⁱ	0.96 (3)	2.58 (4)	3.475 (5)	155 (3)

Symmetry codes: (ii) $-x+1, y+1/2, -z+1/2$; (iii) $-x, -y+1, -z$.