

## 6,11-Dihydroxynaphthacene-5,12-dione

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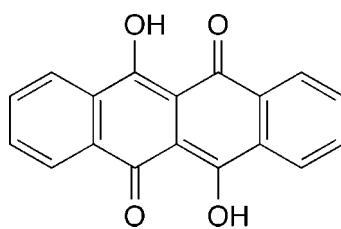
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Key indicators: single-crystal X-ray study;  $T = 173$  K; mean  $\sigma(C-C) = 0.009$  Å;  $R$  factor = 0.114;  $wR$  factor = 0.354; data-to-parameter ratio = 13.0.

The molecule of the title compound,  $C_{18}H_{10}O_4$ , is centrosymmetric and planar. A long phenolic O–H bond is observed [1.19 (9) Å], which is involved in an intramolecular hydrogen bond between the phenolic and quinonoid O atoms. The molecules pack in a herringbone pattern and are linked to each other via intermolecular C–H···O hydrogen bonds (2.73–2.77 Å).

### Related literature

The crystal structure of the rhenium complex including the title compound has been reported (Sathyendiran *et al.*, 2007). For studies of naphthazarin (5,8-dihydroxy-1,4-naphthoquinone), see: Fehlmann & Nigli (1965); Cradwick & Hall (1971); Herbstein *et al.* (1985); Rubio *et al.* (1985); Sarkhel *et al.* (2001); Savko *et al.* (2007). For background on intramolecular hydrogen bonds, see: Gilli *et al.* (1989); Bertolasi *et al.* (1991); Gilli *et al.* (1993); Steiner & Saenger (1994). For background on intermolecular hydrogen bonds, see: Taylor & Kennard (1982); Jagarlapudi & Desiraju (1987); Biradha *et al.* (1993); Batchelor *et al.* (2000). For background on resonance structures, see: Cradwick & Hall (1971); Shiau *et al.* (1980).



### Experimental

#### Crystal data

$C_{18}H_{10}O_4$   
 $M_r = 290.26$   
Monoclinic,  $P2_1/c$

$a = 8.85$  (2) Å  
 $b = 3.750$  (8) Å  
 $c = 18.74$  (4) Å

$\beta = 94.55$  (3)°  
 $V = 620$  (2) Å<sup>3</sup>  
 $Z = 2$   
Mo  $K\alpha$  radiation

$\mu = 0.11$  mm<sup>-1</sup>  
 $T = 173$  (2) K  
 $0.29 \times 0.07 \times 0.03$  mm

#### Data collection

Rigaku Mercury CCD  
diffractometer  
Absorption correction: none  
5026 measured reflections

1347 independent reflections  
668 reflections with  $I > 2\sigma(I)$   
 $R_{\text{int}} = 0.087$

#### Refinement

$R[F^2 > 2\sigma(F^2)] = 0.114$   
 $wR(F^2) = 0.354$   
 $S = 0.99$   
1347 reflections  
104 parameters

H atoms treated by a mixture of  
independent and constrained  
refinement  
 $\Delta\rho_{\text{max}} = 0.52$  e Å<sup>-3</sup>  
 $\Delta\rho_{\text{min}} = -0.32$  e Å<sup>-3</sup>

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

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
O2–H2···O1	1.19 (9)	1.56 (9)	2.516 (7)	132 (6)
C6–H6···O1 <sup>i</sup>	0.95	2.77	3.449 (9)	129
C6–H6···O1 <sup>ii</sup>	0.95	2.74	3.382 (9)	126
C3–H3···O2 <sup>iii</sup>	0.95	2.75	3.378 (10)	124
C4–H4···O2 <sup>iii</sup>	0.95	2.73	3.371 (9)	126

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

Data collection: *CrystalClear* (Molecular Structure Corporation & Rigaku, 2001); cell refinement: *CrystalClear*; data reduction: *TEXSAN* (Molecular Structure Corporation & Rigaku, 2000); program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *PLATON* (Spek, 2003); software used to prepare material for publication: *SHELXL97*.

This work was supported by Grants-in-Aid (Nos. 17750037 and 19550034) from the Ministry of Education, Culture, Sports, Science and Technology, Japan. The authors thank the Instrument Center of the Institute for Molecular Science for the X-ray structure analysis.

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: GW2022).

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# supporting information

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## 6,11-Dihydroxynaphthacene-5,12-dione

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### S1. Comment

Organic molecules containing naphthazarin (5,8-dihydroxy-1,4-naphthoquinone) skeleton have been the attractive subject of structural investigations (Fehlmann & Nigli, 1965; Cradwick & Hall, 1971; Herbstein *et al.*, 1985; Rubio *et al.*, 1985; Sarkhel *et al.*, 2001; Savko *et al.*, 2007). The title compound, (I) (Fig. 1), is  $\pi$ -extended naphthazarin and its molecular and crystal structures are described here.

The molecule of (I) is centrosymmetric and planar, with an r.m.s deviation of 0.0098 Å from the least-squares plane for the fitted non-H atoms. The molecular structure is similar to that of naphthazarin. The symmetry of the carbon skeleton of (I) is close to  $D_{2h}$  (Table 1). The long phenolic O2—H2 bond [1.19 (9) Å] constitutes an intramolecular hydrogen bond between the phenolic O2 and quinonoid O1 atoms (Gilli *et al.*, 1989; Bertolasi *et al.*, 1991; Gilli *et al.*, 1993; Steiner & Saenger, 1994). These facts imply that the structure of (I) can be interpreted not as 1,4- nor 1,5-quinone but as resonance between two zwitterion contributors, as shown in Scheme 1 (Cradwick & Hall, 1971; Shiau *et al.*, 1980).

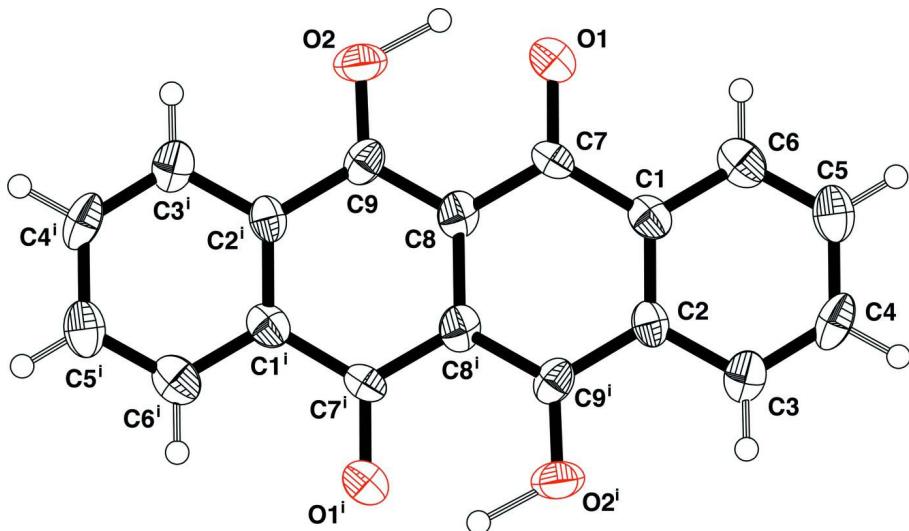
In the crystal structure, the molecules form a herringbone-type stacking along the *b* axis, where the distance between the molecular planes is 3.42 Å (Fig. 2). The packing mode is similar to that characteristic of aromatic hydrocarbon atoms. The intermolecular C—H $\cdots$ O hydrogen bonds (Taylor & Kennard, 1982; Jagarlapudi & Desiraju, 1987; Biradha *et al.*, 1993; Batchelor *et al.*, 2000) are found between the herringbone-type stackings (Table 2). Similar C—H $\cdots$ O hydrogen bonds (2.54–3.01 Å) were observed in the crystal structure of naphthazarin (Cradwick & Hall, 1971).

### S2. Experimental

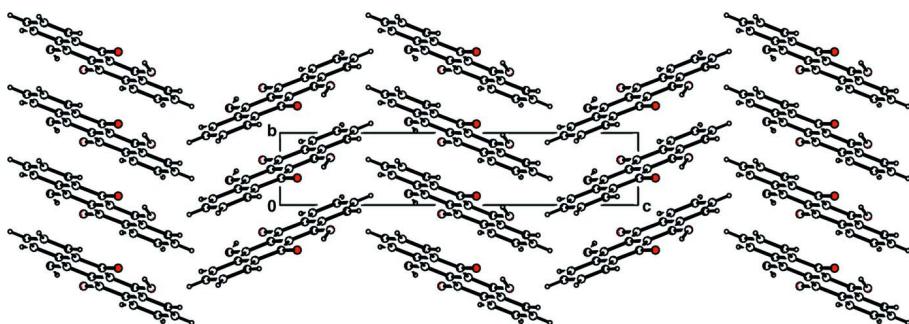
The title compound (I) was commercially available. Red crystals of (I) suitable for X-ray analysis were grown from a chloroform solution.

### S3. Refinement

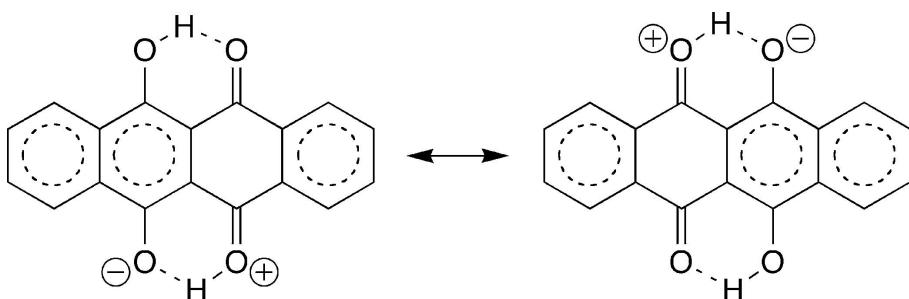
The H atom bonded to the phenolic O atom was located in a difference map and refined isotropically. Other H atoms were positioned geometrically refined using a riding model with C—H = 0.95 Å and with  $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$ .

**Figure 1**

The molecular structure of (I), with atom labels and 50% probability displacement ellipsoids for non-H atoms and H atoms are shown as small spheres of arbitrary radii [symmetry code: (i)  $-x, -y, -z + 1$ ].

**Figure 2**

The packing diagram of (I), viewed along the  $a$  axis.

**Figure 3**

The resonance structure of (I).

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#### Crystal data

$C_{18}H_{10}O_4$   
 $M_r = 290.26$

Monoclinic,  $P2_1/c$   
Hall symbol: -P 2ybc

$a = 8.85 (2)$  Å  
 $b = 3.750 (8)$  Å  
 $c = 18.74 (4)$  Å  
 $\beta = 94.55 (3)^\circ$   
 $V = 620 (2)$  Å<sup>3</sup>  
 $Z = 2$   
 $F(000) = 300$   
 $D_x = 1.555$  Mg m<sup>-3</sup>

Mo  $K\alpha$  radiation,  $\lambda = 0.71070$  Å  
Cell parameters from 928 reflections  
 $\theta = 3.1\text{--}27.5^\circ$   
 $\mu = 0.11$  mm<sup>-1</sup>  
 $T = 173$  K  
Prism, red  
 $0.29 \times 0.07 \times 0.03$  mm

#### Data collection

Rigaku Mercury CCD  
diffractometer  
Radiation source: Rotating Anode  
Graphite Monochromator monochromator  
Detector resolution: 14.6199 pixels mm<sup>-1</sup>  
 $\varphi$  and  $\omega$  scans  
5026 measured reflections

1347 independent reflections  
668 reflections with  $I > 2\sigma(I)$   
 $R_{\text{int}} = 0.087$   
 $\theta_{\text{max}} = 27.5^\circ, \theta_{\text{min}} = 3.1^\circ$   
 $h = -9 \rightarrow 11$   
 $k = -4 \rightarrow 3$   
 $l = -19 \rightarrow 24$

#### Refinement

Refinement on  $F^2$   
Least-squares matrix: full  
 $R[F^2 > 2\sigma(F^2)] = 0.114$   
 $wR(F^2) = 0.354$   
 $S = 0.99$   
1347 reflections  
104 parameters  
0 restraints  
Primary atom site location: structure-invariant  
direct methods

Secondary atom site location: difference Fourier  
map  
Hydrogen site location: inferred from  
neighbouring sites  
H atoms treated by a mixture of independent  
and constrained refinement  
 $w = 1/[\sigma^2(F_o^2) + (0.1532P)^2 + 1.2875P]$   
where  $P = (F_o^2 + 2F_c^2)/3$   
 $(\Delta/\sigma)_{\text{max}} = 0.053$   
 $\Delta\rho_{\text{max}} = 0.52$  e Å<sup>-3</sup>  
 $\Delta\rho_{\text{min}} = -0.32$  e Å<sup>-3</sup>

#### 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 (Å<sup>2</sup>)

	$x$	$y$	$z$	$U_{\text{iso}}^*/U_{\text{eq}}$
O1	0.3137 (5)	0.1246 (12)	0.4518 (2)	0.0438 (12)
O2	0.1298 (5)	-0.1563 (12)	0.3591 (2)	0.0426 (12)
H2	0.248 (10)	-0.03 (3)	0.378 (5)	0.09 (3)*
C1	0.2384 (6)	0.2682 (14)	0.5696 (3)	0.0317 (13)
C2	0.1249 (6)	0.2760 (13)	0.6171 (3)	0.0290 (12)
C3	0.1583 (7)	0.4134 (15)	0.6871 (3)	0.0363 (14)
H3	0.0823	0.4198	0.7201	0.044*
C4	0.3022 (7)	0.5375 (15)	0.7068 (3)	0.0381 (15)
H4	0.3247	0.6318	0.7535	0.046*

C5	0.4162 (7)	0.5264 (15)	0.6586 (3)	0.0403 (15)
H5	0.5151	0.6117	0.6728	0.048*
C6	0.3839 (7)	0.3917 (14)	0.5910 (3)	0.0371 (14)
H6	0.4611	0.3823	0.5586	0.045*
C7	0.2079 (6)	0.1277 (14)	0.4965 (3)	0.0294 (12)
C8	0.0599 (6)	-0.0051 (14)	0.4744 (3)	0.0312 (13)
C9	0.0278 (7)	-0.1452 (13)	0.4050 (3)	0.0308 (13)

Atomic displacement parameters ( $\text{\AA}^2$ )

	$U^{11}$	$U^{22}$	$U^{33}$	$U^{12}$	$U^{13}$	$U^{23}$
O1	0.034 (2)	0.052 (3)	0.046 (3)	-0.011 (2)	0.0070 (18)	-0.004 (2)
O2	0.042 (3)	0.048 (3)	0.041 (2)	0.000 (2)	0.0186 (19)	-0.0051 (19)
C1	0.027 (3)	0.025 (3)	0.043 (3)	0.003 (2)	0.002 (2)	-0.001 (2)
C2	0.028 (3)	0.024 (3)	0.034 (3)	0.000 (2)	-0.005 (2)	0.001 (2)
C3	0.049 (4)	0.028 (3)	0.031 (3)	-0.005 (3)	-0.003 (2)	0.004 (2)
C4	0.063 (4)	0.027 (3)	0.023 (3)	-0.001 (3)	-0.008 (3)	0.000 (2)
C5	0.042 (3)	0.031 (3)	0.045 (3)	-0.001 (3)	-0.011 (3)	0.002 (3)
C6	0.038 (3)	0.022 (3)	0.052 (4)	-0.004 (2)	0.004 (3)	0.005 (2)
C7	0.021 (3)	0.030 (3)	0.037 (3)	-0.001 (2)	0.005 (2)	0.002 (2)
C8	0.036 (3)	0.031 (3)	0.025 (3)	0.001 (2)	-0.006 (2)	0.012 (2)
C9	0.044 (3)	0.021 (2)	0.028 (3)	-0.002 (2)	0.002 (2)	-0.002 (2)

Geometric parameters ( $\text{\AA}$ ,  $^\circ$ )

O1—C7	1.304 (6)	C4—C5	1.407 (9)
O2—C9	1.296 (7)	C4—H4	0.9500
O2—H2	1.19 (9)	C5—C6	1.373 (8)
C1—C2	1.395 (8)	C5—H5	0.9500
C1—C6	1.398 (8)	C6—H6	0.9500
C1—C7	1.472 (8)	C7—C8	1.432 (8)
C2—C3	1.418 (8)	C8—C9	1.410 (7)
C2—C9 <sup>i</sup>	1.467 (8)	C8—C8 <sup>i</sup>	1.486 (11)
C3—C4	1.378 (9)	C9—C2 <sup>i</sup>	1.467 (8)
C3—H3	0.9500		
C7—O1—H2	110 (3)	C6—C5—H5	120.1
C9—O2—H2	115 (4)	C4—C5—H5	120.1
C2—C1—C6	120.2 (5)	C5—C6—C1	120.4 (6)
C2—C1—C7	120.9 (5)	C5—C6—H6	119.8
C6—C1—C7	118.9 (5)	C1—C6—H6	119.8
C1—C2—C3	119.4 (5)	O1—C7—C8	119.9 (5)
C1—C2—C9 <sup>i</sup>	120.4 (5)	O1—C7—C1	120.8 (5)
C3—C2—C9 <sup>i</sup>	120.2 (5)	C8—C7—C1	119.3 (5)
C2—C3—C4	119.4 (5)	C9—C8—C7	120.6 (5)
C2—C3—H3	120.3	C9—C8—C8 <sup>i</sup>	120.2 (6)
C4—C3—H3	120.3	C7—C8—C8 <sup>i</sup>	119.2 (6)
C3—C4—C5	120.8 (5)	O2—C9—C8	121.6 (5)

C3—C4—H4	119.6	O2—C9—C2 <sup>i</sup>	118.4 (5)
C5—C4—H4	119.6	C8—C9—C2 <sup>i</sup>	119.9 (5)
C6—C5—C4	119.7 (6)		
C6—C1—C2—C3	0.6 (8)	C6—C1—C7—O1	-1.4 (8)
C7—C1—C2—C3	-180.0 (5)	C2—C1—C7—C8	-0.5 (8)
C6—C1—C2—C9 <sup>i</sup>	-179.7 (5)	C6—C1—C7—C8	178.9 (5)
C7—C1—C2—C9 <sup>i</sup>	-0.3 (8)	O1—C7—C8—C9	1.1 (7)
C1—C2—C3—C4	0.2 (7)	C1—C7—C8—C9	-179.3 (5)
C9 <sup>i</sup> —C2—C3—C4	-179.5 (5)	O1—C7—C8—C8 <sup>i</sup>	-179.1 (6)
C2—C3—C4—C5	-0.6 (8)	C1—C7—C8—C8 <sup>i</sup>	0.6 (8)
C3—C4—C5—C6	0.3 (8)	C7—C8—C9—O2	-0.5 (8)
C4—C5—C6—C1	0.5 (8)	C8 <sup>i</sup> —C8—C9—O2	179.7 (6)
C2—C1—C6—C5	-0.9 (8)	C7—C8—C9—C2 <sup>i</sup>	-179.2 (5)
C7—C1—C6—C5	179.6 (5)	C8 <sup>i</sup> —C8—C9—C2 <sup>i</sup>	1.0 (8)
C2—C1—C7—O1	179.2 (5)		

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

#### Hydrogen-bond geometry ( $\text{\AA}$ , °)

D—H···A	D—H	H···A	D···A	D—H···A
O2—H2···O1	1.19 (9)	1.56 (9)	2.516 (7)	132 (6)
C6—H6···O1 <sup>ii</sup>	0.95	2.77	3.449 (9)	129
C6—H6···O1 <sup>iii</sup>	0.95	2.74	3.382 (9)	126
C3—H3···O2 <sup>iv</sup>	0.95	2.75	3.378 (10)	124
C4—H4···O2 <sup>iv</sup>	0.95	2.73	3.371 (9)	126

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