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4'-Hydroxybiphenyl-4-carboxylic acid

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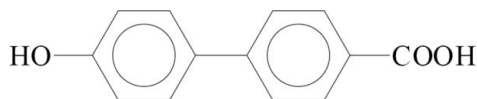
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 Key indicators: single-crystal X-ray study; $T = 293$ K; mean $\sigma(\text{C}-\text{C}) = 0.005$ Å; disorder in main residue; R factor = 0.059; wR factor = 0.184; data-to-parameter ratio = 11.2.

The title compound, $\text{C}_{13}\text{H}_{10}\text{O}_3$, has potential oxygen donor and acceptor sites. Intermolecular hydrogen bonding between neighboring carboxylate groups leads to the formation of hydrogen-bonded dimers [graph-set motif $R_2^2(8)$]. A second hydrogen-bonding interaction between the hydroxy groups generates a chain and extends the structure into a lamellar layer. One of the benzene rings is disordered over two positions with an occupancy ratio of 0.57 (2):0.43 (2).

Related literature

For related literature, see: Bernstein *et al.* (1995); Datta & Pati (2006); Zwier *et al.* (1996).



Experimental

Crystal data

$\text{C}_{13}\text{H}_{10}\text{O}_3$
 $M_r = 214.21$
 Monoclinic, $P2_1/n$
 $a = 8.6500$ (7) Å

$b = 5.5077$ (5) Å
 $c = 20.9655$ (18) Å
 $\beta = 94.145$ (3)°
 $V = 996.22$ (15) Å³

$Z = 4$
 Mo $K\alpha$ radiation
 $\mu = 0.10$ mm⁻¹

$T = 293$ (2) K
 $0.21 \times 0.20 \times 0.16$ mm

Data collection

Bruker APEXII area-detector diffractometer
 Absorption correction: none
 6310 measured reflections

1800 independent reflections
 854 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.063$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.058$
 $wR(F^2) = 0.183$
 $S = 1.01$
 1800 reflections
 160 parameters

24 restraints
 H-atom parameters constrained
 $\Delta\rho_{\text{max}} = 0.18$ e Å⁻³
 $\Delta\rho_{\text{min}} = -0.19$ e Å⁻³

Table 1

Hydrogen-bond geometry (Å, °).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
$\text{O2}-\text{H2}\cdots\text{O1}^{\text{i}}$	0.82	1.82	2.624 (3)	168
$\text{O3}-\text{H3A}\cdots\text{O3}^{\text{ii}}$	0.82	2.20	3.0041 (18)	168

Symmetry codes: (i) $-x + 3, -y, -z$; (ii) $-x + \frac{1}{2}, y - \frac{1}{2}, -z + \frac{1}{2}$.

Data collection: *APEX2* (Bruker, 2004); cell refinement: *APEX2*; data reduction: *APEX2*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *XP in SHELXTL* (Sheldrick, 2008); software used to prepare material for publication: *SHELXTL*.

The author thanks South China Normal University for supporting this study.

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: ZL2111).

References

- Bernstein, J., Davis, R. E., Shimon, L. & Chang, N.-L. (1995). *Angew. Chem. Int. Ed.* **34**, 1555–1573.
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supporting information

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4'-Hydroxybiphenyl-4-carboxylic acid

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S1. Comment

Hydrogen-bonding interactions between ligands are specific and directional. When present in metal complexes they usually do not rely on the properties of the metal ions, but they play an important role in the overall structures and functions of the complexes and the way in which they pack in the solid state (Zwier *et al.*, 1996; Datta & Pati, 2006). In this context we report here the crystal structure of the title compound, (I).

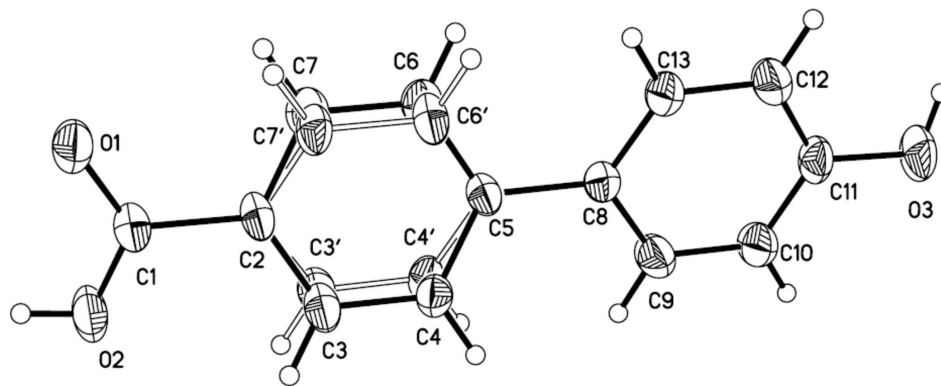
The molecular structure of (I) is shown in Fig. 1. The C—O and C—C bond distances show no remarkable features. The title molecular structure acts as both a hydrogen bonding donor and acceptor, forming dimers with neighboring molecules through O—H \cdots O hydrogen bonding with a $R^2_2(8)$ graph set motif (Bernstein *et al.*, 1995). A second hydrogen bonding interaction by the hydroxyl groups forms a chain and extends the structure into a lamellar layer (Table 1, Fig. 2).

S2. Experimental

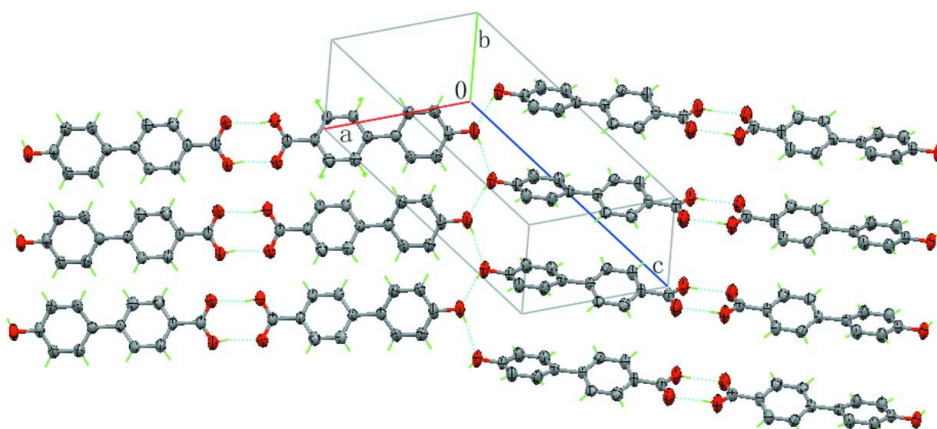
4-Hydroxyl-biphenyl-4'-carboxylic acid was dissolved in a hot ethanol-water solution (1:1; v/v) with stirring. Colorless single crystals suitable for X-ray diffraction were obtained at room temperature by slow evaporation of the solvent over a period of several days.

S3. Refinement

In the initial refinement with disorder not taken into account one of phenyl rings showed significantly elongated thermal ellipsoids indicating disorder, the dihedral angle between two phenyl rings is 5.66 (2) °, and the adjacent distances of C—H \cdots C—H interactions in the biphenylene are 2.044 (1) and 2.077 (1) Å, respectively, thus leading to a static repulsion between two phenyl rings, and the phenyl ring was thus refined as being disordered over two positions. The occupancy ratio refined to 0.57 (2) to 0.43 (2). The ads of the disordered atoms were restrained to be close to isotropic and those of equivalent atoms were set to be identical. Carbon-bound, hydroxyl and carboxylate group H atoms were placed at calculated positions and were treated as riding on their parent C or O atoms with C—H = 0.93 Å, with $U_{\text{iso}}(\text{H}) = 1.2 U_{\text{eq}}(\text{C})$; O—H = 0.82 Å and with $U_{\text{iso}}(\text{H}) = 1.5 U_{\text{eq}}(\text{O})$.

**Figure 1**

The structure of (I), showing the atom-numbering scheme and displacement ellipsoids drawn at the 30% probability level.

**Figure 2**

A layer view of (I).

4'-Hydroxybiphenyl-4-carboxylic acid

Crystal data

$C_{13}H_{10}O_3$

$M_r = 214.21$

Monoclinic, $P2_1/n$

Hall symbol: $-P 2_1/n$

$a = 8.6500 (7) \text{ \AA}$

$b = 5.5077 (5) \text{ \AA}$

$c = 20.9655 (18) \text{ \AA}$

$\beta = 94.145 (3)^\circ$

$V = 996.22 (15) \text{ \AA}^3$

$Z = 4$

$F(000) = 448$

$D_x = 1.428 \text{ Mg m}^{-3}$

Mo $K\alpha$ radiation, $\lambda = 0.71073 \text{ \AA}$

Cell parameters from 1560 reflections

$\theta = 1.4\text{--}28.0^\circ$

$\mu = 0.10 \text{ mm}^{-1}$

$T = 293 \text{ K}$

Plate, colorless

$0.21 \times 0.20 \times 0.16 \text{ mm}$

Data collection

Bruker APEXII area-detector

diffractometer

Radiation source: fine-focus sealed tube

Graphite monochromator

φ and ω scans

6310 measured reflections

1800 independent reflections

854 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.063$

$\theta_{\text{max}} = 25.2^\circ$, $\theta_{\text{min}} = 2.0^\circ$

$h = -10 \rightarrow 10$
 $k = -5 \rightarrow 6$

$l = -24 \rightarrow 25$

Refinement

Refinement on F^2
 Least-squares matrix: full
 $R[F^2 > 2\sigma(F^2)] = 0.058$
 $wR(F^2) = 0.183$
 $S = 1.01$
 1800 reflections
 160 parameters
 24 restraints
 Primary atom site location: structure-invariant
 direct methods

Secondary atom site location: difference Fourier
 map
 Hydrogen site location: inferred from
 neighbouring sites
 H-atom parameters constrained
 $w = 1/[\sigma^2(F_o^2) + (0.0772P)^2]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} < 0.001$
 $\Delta\rho_{\max} = 0.18 \text{ e } \text{\AA}^{-3}$
 $\Delta\rho_{\min} = -0.19 \text{ e } \text{\AA}^{-3}$

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	$U_{\text{iso}}^*/U_{\text{eq}}$	Occ. (<1)
C1	1.3074 (3)	0.0387 (7)	0.03781 (15)	0.0499 (9)	
C8	0.7212 (3)	0.1508 (6)	0.15024 (14)	0.0448 (8)	
C9	0.6208 (3)	0.3440 (6)	0.13452 (16)	0.0578 (10)	
H9	0.6499	0.4597	0.1054	0.069*	
C10	0.4806 (3)	0.3688 (7)	0.16071 (17)	0.0593 (10)	
H10	0.4167	0.4999	0.1493	0.071*	
C11	0.4352 (3)	0.2019 (7)	0.20337 (15)	0.0525 (9)	
C12	0.5302 (4)	0.0092 (7)	0.22074 (16)	0.0596 (10)	
H12	0.5001	-0.1052	0.2500	0.072*	
C13	0.6703 (4)	-0.0119 (6)	0.19414 (15)	0.0545 (9)	
H13	0.7339	-0.1424	0.2064	0.065*	
O1	1.3852 (2)	-0.1522 (5)	0.05016 (11)	0.0704 (8)	
O2	1.3483 (2)	0.2037 (5)	0.00069 (12)	0.0696 (8)	
H2	1.4342	0.1717	-0.0112	0.104*	
O3	0.2936 (3)	0.2362 (5)	0.22900 (12)	0.0720 (8)	
H3A	0.2672	0.1099	0.2458	0.108*	
C2	1.15866 (19)	0.0702 (5)	0.06838 (10)	0.0480 (8)	0.43 (2)
C3	1.0867 (8)	0.2959 (6)	0.0682 (6)	0.052 (3)	0.43 (2)
H3	1.1327	0.4288	0.0498	0.063*	0.43 (2)
C4	0.9458 (8)	0.3228 (6)	0.0954 (6)	0.045 (2)	0.43 (2)
H4	0.8976	0.4738	0.0952	0.054*	0.43 (2)
C5	0.8770 (2)	0.1241 (4)	0.12279 (11)	0.0441 (8)	0.43 (2)

C6	0.9490 (7)	-0.1015 (7)	0.1230 (5)	0.045 (3)	0.43 (2)
H6	0.9029	-0.2345	0.1413	0.054*	0.43 (2)
C7	1.0898 (8)	-0.1285 (7)	0.0958 (6)	0.052 (3)	0.43 (2)
H7	1.1380	-0.2795	0.0959	0.062*	0.43 (2)
C2'	1.15879 (19)	0.0709 (5)	0.06822 (10)	0.0480 (8)	0.57 (2)
C3'	1.0571 (7)	0.2579 (15)	0.0469 (5)	0.052 (2)	0.57 (2)
H3'	1.0845	0.3647	0.0153	0.063*	0.57 (2)
C4'	0.9160 (7)	0.2818 (15)	0.0734 (5)	0.049 (2)	0.57 (2)
H4'	0.8485	0.4045	0.0588	0.059*	0.57 (2)
C5'	0.8725 (2)	0.1234 (5)	0.12205 (11)	0.0441 (8)	0.57 (2)
C6'	0.9755 (6)	-0.0581 (16)	0.1431 (4)	0.048 (2)	0.57 (2)
H6'	0.9505	-0.1621	0.1758	0.057*	0.57 (2)
C7'	1.1161 (6)	-0.0845 (15)	0.1155 (4)	0.047 (2)	0.57 (2)
H7'	1.1827	-0.2097	0.1291	0.056*	0.57 (2)

Atomic displacement parameters (Å²)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
C1	0.0396 (18)	0.061 (2)	0.051 (2)	0.0010 (18)	0.0134 (16)	0.0005 (19)
C8	0.0388 (17)	0.044 (2)	0.0527 (18)	-0.0001 (16)	0.0113 (15)	-0.0022 (17)
C9	0.0445 (19)	0.058 (2)	0.073 (2)	0.0005 (18)	0.0159 (17)	0.011 (2)
C10	0.0418 (19)	0.057 (2)	0.081 (2)	0.0079 (18)	0.0170 (18)	0.008 (2)
C11	0.0338 (17)	0.062 (3)	0.064 (2)	-0.0032 (17)	0.0160 (15)	-0.0135 (19)
C12	0.049 (2)	0.061 (3)	0.071 (2)	0.0015 (18)	0.0214 (18)	0.0075 (19)
C13	0.0448 (19)	0.051 (2)	0.069 (2)	0.0088 (17)	0.0183 (17)	0.0053 (18)
O1	0.0529 (15)	0.0705 (18)	0.0910 (18)	0.0171 (13)	0.0281 (13)	0.0188 (15)
O2	0.0460 (14)	0.083 (2)	0.0837 (18)	0.0127 (13)	0.0307 (13)	0.0194 (15)
O3	0.0418 (13)	0.090 (2)	0.0882 (18)	0.0033 (13)	0.0308 (12)	-0.0057 (16)
C2	0.0334 (17)	0.062 (2)	0.050 (2)	-0.0008 (17)	0.0144 (15)	0.0031 (18)
C3	0.040 (5)	0.065 (7)	0.054 (5)	0.002 (5)	0.015 (4)	0.006 (5)
C4	0.046 (5)	0.047 (6)	0.043 (5)	0.005 (4)	0.012 (4)	0.000 (4)
C5	0.0367 (17)	0.049 (2)	0.0476 (19)	-0.0041 (16)	0.0129 (14)	0.0004 (17)
C6	0.045 (5)	0.052 (5)	0.039 (5)	-0.002 (4)	0.002 (4)	-0.008 (4)
C7	0.041 (5)	0.061 (7)	0.052 (5)	0.003 (4)	0.001 (4)	-0.001 (5)
C2'	0.0334 (17)	0.062 (2)	0.050 (2)	-0.0008 (17)	0.0144 (15)	0.0031 (18)
C3'	0.044 (4)	0.057 (5)	0.058 (4)	0.004 (3)	0.014 (4)	0.008 (4)
C4'	0.042 (4)	0.049 (4)	0.057 (4)	0.012 (3)	0.014 (3)	0.000 (4)
C5'	0.0367 (17)	0.049 (2)	0.0476 (19)	-0.0041 (16)	0.0129 (14)	0.0004 (17)
C6'	0.036 (4)	0.068 (5)	0.039 (4)	0.007 (3)	0.010 (3)	0.009 (4)
C7'	0.029 (3)	0.063 (5)	0.049 (4)	0.011 (3)	0.005 (3)	0.007 (4)

Geometric parameters (Å, °)

C1—O2	1.264 (4)	C3—C4	1.3900
C1—O1	1.265 (4)	C3—H3	0.9300
C1—C2	1.488 (4)	C4—C5	1.3900
C8—C13	1.379 (4)	C4—H4	0.9300
C8—C9	1.398 (4)	C5—C6	1.3900

C8—C5	1.510 (3)	C6—C7	1.3900
C9—C10	1.374 (4)	C6—H6	0.9300
C9—H9	0.9300	C7—H7	0.9300
C10—C11	1.360 (5)	C2'—C7'	1.3793
C10—H10	0.9300	C2'—C3'	1.4061
C11—C12	1.375 (4)	C3'—C4'	1.3820
C11—O3	1.386 (4)	C3'—H3'	0.9300
C12—C13	1.375 (4)	C4'—C5'	1.4147
C12—H12	0.9300	C4'—H4'	0.9300
C13—H13	0.9300	C5'—C6'	1.3903
O2—H2	0.8200	C6'—C7'	1.3923
O3—H3A	0.8200	C6'—H6'	0.9300
C2—C3	1.3900	C7'—H7'	0.9300
C2—C7	1.3900		
O2—C1—O1	123.7 (3)	C3—C4—C5	120.0
O2—C1—C2	118.1 (3)	C3—C4—H4	120.0
O1—C1—C2	118.2 (3)	C5—C4—H4	120.0
C13—C8—C9	115.4 (3)	C6—C5—C4	120.0
C13—C8—C5	121.9 (3)	C6—C5—C8	119.8 (2)
C9—C8—C5	122.7 (3)	C4—C5—C8	120.1 (2)
C10—C9—C8	122.2 (3)	C5—C6—C7	120.0
C10—C9—H9	118.9	C5—C6—H6	120.0
C8—C9—H9	118.9	C7—C6—H6	120.0
C11—C10—C9	120.1 (3)	C6—C7—C2	120.0
C11—C10—H10	120.0	C6—C7—H7	120.0
C9—C10—H10	120.0	C2—C7—H7	120.0
C10—C11—C12	120.1 (3)	C7'—C2'—C3'	119.2
C10—C11—O3	117.9 (3)	C4'—C3'—C2'	119.5
C12—C11—O3	122.0 (3)	C4'—C3'—H3'	120.2
C11—C12—C13	119.0 (3)	C2'—C3'—H3'	120.2
C11—C12—H12	120.5	C3'—C4'—C5'	121.3
C13—C12—H12	120.5	C3'—C4'—H4'	119.4
C12—C13—C8	123.3 (3)	C5'—C4'—H4'	119.4
C12—C13—H13	118.4	C6'—C5'—C4'	118.4
C8—C13—H13	118.4	C5'—C6'—C7'	120.1
C3—C2—C7	120.0	C5'—C6'—H6'	120.0
C3—C2—C1	120.3 (2)	C7'—C6'—H6'	120.0
C7—C2—C1	119.7 (2)	C2'—C7'—C6'	121.5
C4—C3—C2	120.0	C2'—C7'—H7'	119.3
C4—C3—H3	120.0	C6'—C7'—H7'	119.3
C2—C3—H3	120.0		

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

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
O2—H2 \cdots O1 ⁱ	0.82	1.82	2.624 (3)	168

O3—H3A···O3 ⁱⁱ	0.82	2.20	3.0041 (18)	168
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Symmetry codes: (i) $-x+3, -y, -z$; (ii) $-x+1/2, y-1/2, -z+1/2$.