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4'-Formylbiphenyl-4-yl acetate

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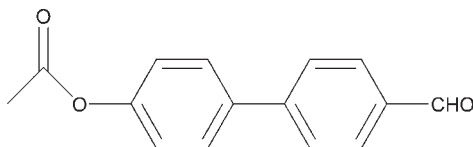
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Key indicators: single-crystal X-ray study; $T = 298$ K; mean $\sigma(\text{C}-\text{C}) = 0.004$ Å;
 R factor = 0.034; wR factor = 0.091; data-to-parameter ratio = 7.9.

In the title compound, $\text{C}_{15}\text{H}_{12}\text{O}_3$, the dihedral angle between the six-membered rings is $30.39(1)^\circ$. The crystal packing is stabilized by intermolecular $\text{C}-\text{H}\cdots\text{O}$ hydrogen bonds.

Related literature

For further synthetic details, see: Chakraborti & Gulhane (2003); Chamontin *et al.* (1999); Steglich & Höfle (1969).



Experimental

Crystal data

 $\text{C}_{15}\text{H}_{12}\text{O}_3$ $M_r = 240.25$ Monoclinic, $P2_1$ $a = 9.250(6)$ Å $b = 7.499(4)$ Å $c = 9.596(6)$ Å $\beta = 113.695(10)^\circ$ $V = 609.5(6)$ Å³ $Z = 2$ Mo $K\alpha$ radiation $\mu = 0.09$ mm⁻¹ $T = 298$ K $0.18 \times 0.15 \times 0.10$ mm

Data collection

Bruker SMART APEXII CCD-
detector diffractometer
Absorption correction: multi-scan
(*SADABS*; Sheldrick, 2004)
 $T_{\min} = 0.984$, $T_{\max} = 0.991$ 3288 measured reflections
1290 independent reflections
1077 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.019$

Refinement

 $R[F^2 > 2\sigma(F^2)] = 0.034$ $wR(F^2) = 0.091$ $S = 1.00$

1290 reflections

164 parameters

1 restraint

H-atom parameters constrained

 $\Delta\rho_{\text{max}} = 0.09$ e Å⁻³ $\Delta\rho_{\text{min}} = -0.12$ e Å⁻³

Table 1

Hydrogen-bond geometry (Å, °).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
$\text{C15}-\text{H15C}\cdots\text{O3}^i$	0.96	2.41	3.372 (5)	177

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

Data collection: *APEX2* (Bruker, 2004); cell refinement: *SAINT* (Bruker, 2004); data reduction: *SAINT*; 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: *SHELXL97*.

This work was supported by the Natural Science Foundation of Ludong University and the Students Research Fund of Ludong University.

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: BG2320).

References

- Bruker (2004). *APEX2* and *SAINT*. Bruker AXS Inc., Madison, Wisconsin, USA.
- Chakraborti, A. K. & Gulhane, R. (2003). *Tetrahedron Lett.* **44**, 3521–3525.
- Chamontin, K., Lokshin, V., Rossollin, V., Samat, A. & Guglielmetti, R. (1999). *Tetrahedron*, **55**, 5821–5830.
- Sheldrick, G. M. (2004). *SADABS*. University of Göttingen, Germany.
- Sheldrick, G. M. (2008). *Acta Cryst.* **A64**, 112–122.
- Steglich, W. & Höfle, G. (1969). *Angew. Chem. Int. Ed. Engl.* **8**, 981–981.

supporting information

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4'-Formylbiphenyl-4-yl acetate

Sheng-guang Xu, Jun-shen Liu, Chun-hua Wang, Chun-nuan Ji and Gang Liu

S1. Comment

4'-formylbiphenyl-4-yl acetate is a derivative of biphenyl that contains two phenyl groups joined by a single covalent bond. However, rotation about the single bond in biphenyl may be sterically hindered and the equilibrium torsional angles in the variously substituted derivatives may be different. In the case of unsubstituted biphenyl, the equilibrium torsional angle is 44.4°. However, in the title compound its value is 30.39 (1)°, because conjugated effects in the molecule are strengthened by π - π conjugation (aldehyde group, oxygen group) as compared with unsubstituted biphenyl, and as result the two phenyl groups tend to be more coplanar.

The molecule of the title compound is illustrated in Fig. 1. The asymmetric unit contains two six-member rings and one acetate group. In the crystal structure there are no classic hydrogen bonds but there are non-classical intermolecular C—H \cdots O hydrogen bonds (viz., C15—H15C \cdots O3[-x,1/2+y,2-z], H \cdots O: 2.41 Å, C—H \cdots O: 177°) which help in forming a three dimensional structure.

S2. Experimental

4'-hydroxy-4-biphenylaldehyde (**02**) was prepared (Chamontin *et al.* 1999), starting from 4'-bromo-4-biphenol (**01**) which is commercially available, using the *N*-formylpiperidine as electrophilic agent. After 4'-hydroxy-4-biphenylaldehyde (**02**) was acetylated with the acetylating agent acetic anhydride, catalyzed by 4-(dimthylamino)pyridine (DMAP), the target material 4'-formylbiphenyl-4-ylacetate (**B₃**) was obtained (Chakraborti *et al.* 2003; Steglich *et al.* 1969).

Product, yielding 53%. ¹H NMR (400MHz, CDCl₃, TMS) δ 10.0939 (s, 1H), δ 7.996–7.976 (d, 2H), δ 7.772–7.752 (d, 2H), δ 7.688–7.666 (d, 2H), δ 7.256–7.235 (d, 2H), 2.374 (s, 3H); ¹³C NMR (CDCl₃) δ 191.79, 169.40, 151.03, 146.28, 137.46, 135.25, 130.29, 128.45, 127.64, 122.19, 21.147; ESI-TOF Eaxct Mass for C₁₅H₁₂O₃ [M+Na]⁺: calcd 263.0679, 264.0713, 265.0738, found 263.0401, 264.0444, 265.0743.

S3. Refinement

All H atoms were positioned in calculated positions, with C—H distances of 0.97 Å, and with $U_{iso}(H) = 1.2$ or $1.5 U_{eq}(C)$. In the absence of significant anomalous effects, Friedel pairs were merged, thus giving rise to a poorer reflections to parameters ratio.

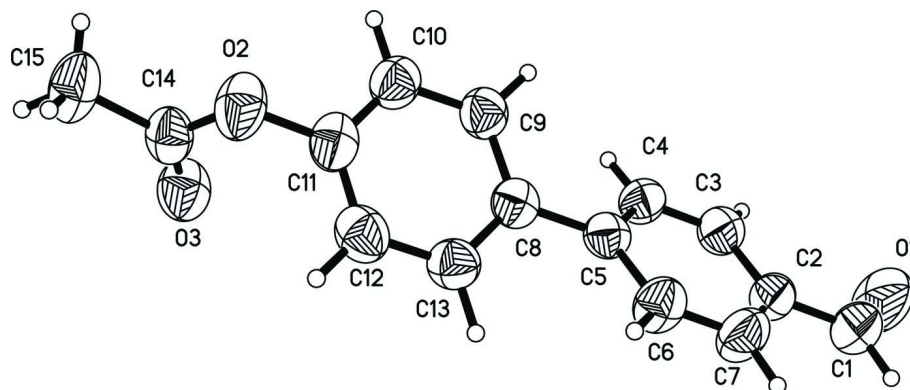


Figure 1

Displacement ellipsoid plot of (I) (50% probability level).

4'-Formylbiphenyl-4-yl acetate

Crystal data

$C_{15}H_{12}O_3$

$M_r = 240.25$

Monoclinic, $P2_1$

Hall symbol: P 2yb

$a = 9.250$ (6) Å

$b = 7.499$ (4) Å

$c = 9.596$ (6) Å

$\beta = 113.695$ (10)°

$V = 609.5$ (6) Å³

$Z = 2$

$F(000) = 252$

$D_x = 1.309$ Mg m⁻³

Mo $K\alpha$ radiation, $\lambda = 0.71073$ Å

Cell parameters from 1236 reflections

$\theta = 2.3$ – 23.9 °

$\mu = 0.09$ mm⁻¹

$T = 298$ K

Block, colourless

$0.18 \times 0.15 \times 0.10$ mm

Data collection

Bruker SMART APEX CCD-detector
diffractometer

Radiation source: fine-focus sealed tube

Graphite monochromator

ω scans

Absorption correction: multi-scan

(*SADABS*; Sheldrick, 2004)

$T_{\min} = 0.984$, $T_{\max} = 0.991$

3288 measured reflections

1290 independent reflections

1077 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.019$

$\theta_{\max} = 26.0$ °, $\theta_{\min} = 2.3$ °

$h = -11 \rightarrow 11$

$k = -7 \rightarrow 9$

$l = -8 \rightarrow 11$

Refinement

Refinement on F^2

Least-squares matrix: full

$R[F^2 > 2\sigma(F^2)] = 0.034$

$wR(F^2) = 0.091$

$S = 1.00$

1290 reflections

164 parameters

1 restraint

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

$w = 1/[\sigma^2(F_o^2) + (0.0452P)^2 + 0.0661P]$

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

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

$\Delta\rho_{\max} = 0.09$ e Å⁻³

$\Delta\rho_{\min} = -0.12$ e Å⁻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	$U_{\text{iso}}^*/U_{\text{eq}}$
O2	-0.0502 (2)	0.8608 (3)	0.7519 (2)	0.0701 (5)
C8	0.2140 (3)	0.8713 (3)	0.4862 (3)	0.0477 (5)
C4	0.2728 (3)	0.7574 (3)	0.2696 (3)	0.0513 (6)
H4	0.1884	0.6789	0.2457	0.062*
C5	0.3071 (2)	0.8734 (3)	0.3918 (2)	0.0473 (5)
C11	0.0414 (3)	0.8599 (3)	0.6654 (3)	0.0560 (6)
C2	0.4874 (3)	0.8714 (4)	0.2161 (3)	0.0554 (6)
C9	0.0550 (3)	0.8248 (3)	0.4255 (3)	0.0542 (6)
H9	0.0052	0.7970	0.3226	0.065*
C13	0.2828 (3)	0.9145 (3)	0.6404 (3)	0.0557 (6)
H13	0.3886	0.9477	0.6841	0.067*
C3	0.3610 (3)	0.7561 (4)	0.1831 (3)	0.0560 (6)
H3	0.3355	0.6771	0.1019	0.067*
C10	-0.0305 (3)	0.8190 (4)	0.5149 (3)	0.0580 (6)
H10	-0.1366	0.7873	0.4725	0.070*
C6	0.4340 (3)	0.9888 (4)	0.4230 (3)	0.0604 (7)
H6	0.4593	1.0690	0.5034	0.073*
C1	0.5827 (3)	0.8732 (5)	0.1248 (4)	0.0741 (8)
H1	0.6653	0.9546	0.1514	0.089*
C12	0.1971 (3)	0.9091 (4)	0.7295 (3)	0.0620 (7)
H12	0.2446	0.9386	0.8321	0.074*
C14	-0.0346 (3)	0.7197 (4)	0.8450 (3)	0.0550 (6)
O3	0.0583 (2)	0.6044 (3)	0.8607 (2)	0.0720 (5)
C7	0.5228 (3)	0.9866 (4)	0.3372 (3)	0.0662 (7)
H7	0.6080	1.0640	0.3614	0.079*
C15	-0.1467 (3)	0.7346 (5)	0.9211 (3)	0.0768 (9)
H15A	-0.1402	0.6290	0.9798	0.115*
H15B	-0.2523	0.7478	0.8455	0.115*
H15C	-0.1196	0.8367	0.9871	0.115*
O1	0.5613 (3)	0.7773 (5)	0.0187 (3)	0.1034 (9)

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
O2	0.0865 (12)	0.0605 (11)	0.0839 (12)	0.0192 (11)	0.0557 (11)	0.0128 (11)
C8	0.0530 (12)	0.0361 (11)	0.0536 (13)	-0.0002 (11)	0.0210 (10)	0.0025 (12)

C4	0.0482 (12)	0.0508 (14)	0.0534 (13)	-0.0096 (11)	0.0187 (10)	-0.0039 (12)
C5	0.0485 (12)	0.0398 (11)	0.0506 (12)	0.0013 (11)	0.0168 (10)	0.0055 (12)
C11	0.0675 (15)	0.0435 (13)	0.0675 (15)	0.0122 (14)	0.0380 (13)	0.0117 (13)
C2	0.0494 (12)	0.0603 (15)	0.0592 (14)	0.0015 (13)	0.0244 (11)	0.0081 (14)
C9	0.0576 (13)	0.0523 (15)	0.0524 (13)	-0.0038 (12)	0.0217 (11)	0.0027 (11)
C13	0.0521 (13)	0.0567 (16)	0.0561 (14)	0.0011 (11)	0.0195 (11)	-0.0020 (12)
C3	0.0566 (13)	0.0593 (15)	0.0522 (13)	-0.0017 (12)	0.0218 (11)	-0.0029 (13)
C10	0.0558 (14)	0.0557 (16)	0.0657 (15)	-0.0019 (12)	0.0276 (12)	0.0042 (13)
C6	0.0643 (15)	0.0554 (15)	0.0647 (15)	-0.0155 (13)	0.0291 (13)	-0.0128 (14)
C1	0.0587 (15)	0.091 (2)	0.0785 (18)	0.0027 (18)	0.0339 (14)	0.012 (2)
C12	0.0697 (16)	0.0643 (18)	0.0523 (13)	0.0084 (13)	0.0249 (13)	-0.0027 (13)
C14	0.0675 (15)	0.0529 (15)	0.0496 (13)	-0.0100 (13)	0.0287 (12)	-0.0101 (12)
O3	0.0852 (12)	0.0607 (11)	0.0787 (13)	0.0094 (11)	0.0420 (11)	0.0143 (10)
C7	0.0590 (15)	0.0645 (17)	0.0779 (18)	-0.0182 (15)	0.0304 (14)	-0.0030 (16)
C15	0.094 (2)	0.078 (2)	0.080 (2)	-0.0186 (17)	0.0573 (18)	-0.0183 (17)
O1	0.0920 (15)	0.143 (2)	0.0995 (16)	-0.0075 (17)	0.0638 (14)	-0.0223 (18)

Geometric parameters (Å, °)

O2—C14	1.355 (3)	C13—C12	1.381 (4)
O2—C11	1.404 (3)	C13—H13	0.9300
C8—C9	1.391 (3)	C3—H3	0.9300
C8—C13	1.394 (3)	C10—H10	0.9300
C8—C5	1.480 (3)	C6—C7	1.378 (4)
C4—C3	1.378 (3)	C6—H6	0.9300
C4—C5	1.391 (3)	C1—O1	1.196 (4)
C4—H4	0.9300	C1—H1	0.9300
C5—C6	1.390 (3)	C12—H12	0.9300
C11—C10	1.360 (4)	C14—O3	1.185 (3)
C11—C12	1.369 (4)	C14—C15	1.493 (3)
C2—C7	1.377 (4)	C7—H7	0.9300
C2—C3	1.385 (4)	C15—H15A	0.9600
C2—C1	1.472 (4)	C15—H15B	0.9600
C9—C10	1.382 (3)	C15—H15C	0.9600
C9—H9	0.9300		
C14—O2—C11	117.2 (2)	C11—C10—C9	119.6 (2)
C9—C8—C13	117.3 (2)	C11—C10—H10	120.2
C9—C8—C5	121.6 (2)	C9—C10—H10	120.2
C13—C8—C5	121.1 (2)	C7—C6—C5	121.2 (2)
C3—C4—C5	121.5 (2)	C7—C6—H6	119.4
C3—C4—H4	119.2	C5—C6—H6	119.4
C5—C4—H4	119.2	O1—C1—C2	124.6 (3)
C6—C5—C4	117.3 (2)	O1—C1—H1	117.7
C6—C5—C8	121.5 (2)	C2—C1—H1	117.7
C4—C5—C8	121.2 (2)	C11—C12—C13	119.5 (2)
C10—C11—C12	121.0 (2)	C11—C12—H12	120.3
C10—C11—O2	118.3 (2)	C13—C12—H12	120.3

C12—C11—O2	120.6 (2)	O3—C14—O2	122.2 (2)
C7—C2—C3	118.5 (2)	O3—C14—C15	127.1 (3)
C7—C2—C1	120.0 (3)	O2—C14—C15	110.7 (2)
C3—C2—C1	121.5 (3)	C6—C7—C2	121.0 (2)
C10—C9—C8	121.4 (2)	C6—C7—H7	119.5
C10—C9—H9	119.3	C2—C7—H7	119.5
C8—C9—H9	119.3	C14—C15—H15A	109.5
C12—C13—C8	121.3 (2)	C14—C15—H15B	109.5
C12—C13—H13	119.4	H15A—C15—H15B	109.5
C8—C13—H13	119.4	C14—C15—H15C	109.5
C4—C3—C2	120.5 (2)	H15A—C15—H15C	109.5
C4—C3—H3	119.8	H15B—C15—H15C	109.5
C2—C3—H3	119.8		
C3—C4—C5—C6	0.2 (4)	C12—C11—C10—C9	0.8 (4)
C3—C4—C5—C8	-178.6 (2)	O2—C11—C10—C9	177.1 (2)
C9—C8—C5—C6	150.4 (3)	C8—C9—C10—C11	0.2 (4)
C13—C8—C5—C6	-30.2 (3)	C4—C5—C6—C7	-0.7 (4)
C9—C8—C5—C4	-30.8 (3)	C8—C5—C6—C7	178.2 (2)
C13—C8—C5—C4	148.6 (2)	C7—C2—C1—O1	179.4 (3)
C14—O2—C11—C10	103.6 (3)	C3—C2—C1—O1	-0.8 (5)
C14—O2—C11—C12	-80.1 (3)	C10—C11—C12—C13	-1.0 (4)
C13—C8—C9—C10	-1.0 (4)	O2—C11—C12—C13	-177.2 (3)
C5—C8—C9—C10	178.5 (2)	C8—C13—C12—C11	0.2 (4)
C9—C8—C13—C12	0.8 (4)	C11—O2—C14—O3	4.2 (4)
C5—C8—C13—C12	-178.6 (2)	C11—O2—C14—C15	-176.4 (2)
C5—C4—C3—C2	0.0 (4)	C5—C6—C7—C2	0.9 (4)
C7—C2—C3—C4	0.2 (4)	C3—C2—C7—C6	-0.7 (4)
C1—C2—C3—C4	-179.6 (3)	C1—C2—C7—C6	179.2 (3)

Hydrogen-bond geometry (Å, °)

<i>D</i> —H \cdots <i>A</i>	<i>D</i> —H	H \cdots <i>A</i>	<i>D</i> \cdots <i>A</i>	<i>D</i> —H \cdots <i>A</i>
C15—H15C \cdots O3 ⁱ	0.96	2.41	3.372 (5)	177

Symmetry code: (i) $-x, y+1/2, -z+2$.