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1,1'-[[1,4-Phenylenebis(methylene)]-bis(oxy)bis(4,1-phenylene)]diethanone

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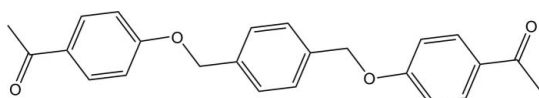
Received 23 October 2011; accepted 31 October 2011

Key indicators: single-crystal X-ray study; $T = 100$ K; mean $\sigma(\text{C}-\text{C}) = 0.002$ Å; R factor = 0.040; wR factor = 0.112; data-to-parameter ratio = 12.9.

The centroid of the central aromatic ring of the title molecule, $\text{C}_{24}\text{H}_{22}\text{O}_4$, is located on an inversion center. The dihedral angle between the central and terminal benzene rings is 75.00 (7)°. In the crystal, molecules are linked through $\text{C}-\text{H}\cdots\text{O}$ hydrogen bonds into chains along $[121]$. The chains are connected into layers *via* $\text{C}-\text{H}\cdots\pi$ interactions.

Related literature

For related structures, see: Al-Mohammed *et al.* (2011); Hu (2010); Tang *et al.* (2008).



Experimental

Crystal data

$\text{C}_{24}\text{H}_{22}\text{O}_4$
 $M_r = 374.42$
Triclinic, $P\bar{1}$
 $a = 8.1286$ (12) Å
 $b = 8.1610$ (7) Å
 $c = 8.4878$ (6) Å
 $\alpha = 116.164$ (5)°
 $\beta = 106.328$ (7)°

$\gamma = 100.196$ (7)°
 $V = 454.41$ (8) Å³
 $Z = 1$
Mo $K\alpha$ radiation
 $\mu = 0.09$ mm⁻¹
 $T = 100$ K
 $0.23 \times 0.19 \times 0.09$ mm

Data collection

Bruker APEXII CCD diffractometer
Absorption correction: multi-scan (*SADABS*; Sheldrick, 1996)
 $T_{\min} = 0.979$, $T_{\max} = 0.992$
2732 measured reflections
1654 independent reflections
1472 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.020$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.040$
 $wR(F^2) = 0.112$
 $S = 1.07$
1654 reflections
128 parameters
H-atom parameters constrained
 $\Delta\rho_{\max} = 0.20$ e Å⁻³
 $\Delta\rho_{\min} = -0.34$ e Å⁻³

Table 1

Hydrogen-bond geometry (Å, °).

Cg1 is the centroid of the $\text{C3}-\text{C8}$ ring.

$\text{D}-\text{H}\cdots\text{A}$	$\text{D}-\text{H}$	$\text{H}\cdots\text{A}$	$\text{D}\cdots\text{A}$	$\text{D}-\text{H}\cdots\text{A}$
$\text{C11}-\text{H11}\cdots\text{O1}^{\text{i}}$	0.95	2.54	3.4362 (18)	158
$\text{C12}-\text{H12}\cdots\text{Cg1}^{\text{ii}}$	0.95	2.61	3.5078 (17)	158

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

Data collection: *APEX2* (Bruker, 2007); cell refinement: *SAINT* (Bruker, 2007); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *X-SEED* (Barbour, 2001); software used to prepare material for publication: *SHELXL97* and *PUBLICIF* (Westrip, 2010).

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: PV2469).

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supporting information

Acta Cryst. (2011). E67, o3164 [https://doi.org/10.1107/S160053681104565X]

1,1'-[1,4-Phenylenebis(methylene)]bis(oxy)bis(4,1-phenylene)diethanone

Nassir N. Al-Mohammed, Yatimah Alias, Zanariah Abdullah and Hamid Khaledi

S1. Comment

We have recently reported the crystal structure of *o*-acetyl isomer of the title compound (Al-Mohammed *et al.*, 2011). Similar to the previous structure, the title molecule shows a centrosymmetric molecular structure with the centroid of the central benzene ring being located on an inversion center. The central and terminal rings make a dihedral angle of 75.00 (7)°. This value is comparable to those observed in the structures of the previously reported isomer and some other similar compounds (Hu, 2010; Tang *et al.*, 2008). In the crystal, the molecules are linked through C—H···O bonds into chains along [1 2 1] direction. The chains are connected into layers *via* C—H··· π interactions (Table 1, Fig. 2).

S2. Experimental

To a suspension of α,α' -dibromo-*p*-xylene (1 g, 3.8 mmol) and potassium carbonate (1.05 g, 7.57 mmol) in dry acetone (25 ml), 4'-hydroxyacetophenone (1.03 g, 7.57 mmole) was added and the mixture was refluxed for 48 hr. The solvent was then evaporated under reduced pressure and the crude material was extracted with dichloromethane (3 x 25 ml). The combined organic layers were washed with water followed by brine and dried over anhydrous sodium sulfate. The solvent was evaporated under vacuum and the formed amorphous solid was re-crystallized from chloroform to obtain colorless crystals of the title compound (m.p. = 435–437 K).

S3. Refinement

Hydrogen atoms were placed at calculated positions and refined in riding mode with C—H distances of 0.95 (aryl), 0.98 (methyl) and 0.99 (methylene) Å, and $U_{\text{iso}}(\text{H})$ set to 1.2 (1.5 for methyl) $U_{\text{eq}}(\text{carrier atoms})$.

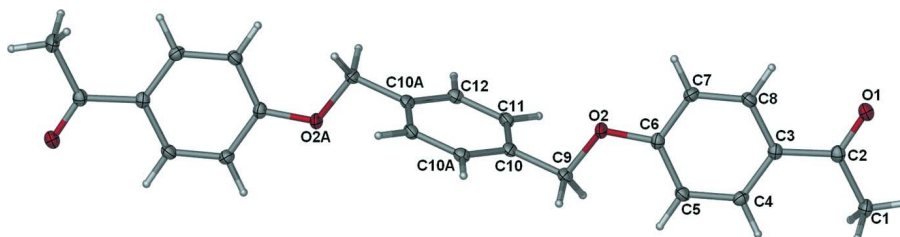


Figure 1

Molecular structure of the title compound with displacement ellipsoids drawn at 50% probability level. Hydrogen atoms are drawn as spheres of arbitrary radius. Symmetry code: A = $-x+1, -y+1, -z+1$.

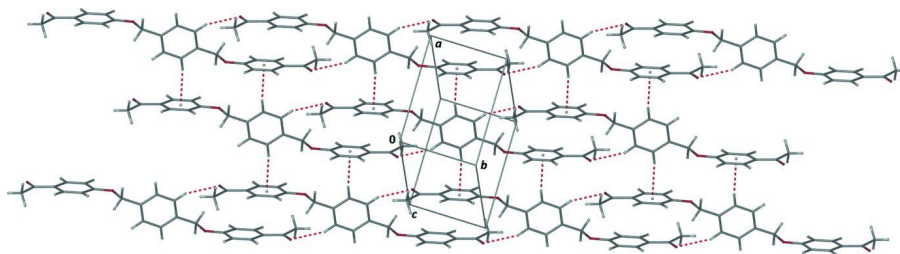


Figure 2

The two-dimensional network formed by C—H...O and C—H... π interactions (dashed lines).

1,1'-[1,4-Phenylenebis(methylene)]bis(oxy)bis(4,1-phenylene)diethanone

Crystal data

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$M_r = 374.42$

Triclinic, $P\bar{1}$

Hall symbol: $-P\ 1$

$a = 8.1286$ (12) Å

$b = 8.1610$ (7) Å

$c = 8.4878$ (6) Å

$\alpha = 116.164$ (5)°

$\beta = 106.328$ (7)°

$\gamma = 100.196$ (7)°

$V = 454.41$ (8) Å³

$Z = 1$

$F(000) = 198$

$D_x = 1.368$ Mg m⁻³

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

Cell parameters from 1506 reflections

$\theta = 2.9$ – 28.8 °

$\mu = 0.09$ mm⁻¹

$T = 100$ K

Block, colorless

$0.23 \times 0.19 \times 0.09$ mm

Data collection

Bruker APEXII CCD
diffractometer

Radiation source: fine-focus sealed tube

Graphite monochromator

φ and ω scans

Absorption correction: multi-scan

(*SADABS*; Sheldrick, 1996)

$T_{\min} = 0.979$, $T_{\max} = 0.992$

2732 measured reflections

1654 independent reflections

1472 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.020$

$\theta_{\max} = 25.5$ °, $\theta_{\min} = 2.8$ °

$h = -9 \rightarrow 9$

$k = -9 \rightarrow 9$

$l = -10 \rightarrow 10$

Refinement

Refinement on F^2

Least-squares matrix: full

$R[F^2 > 2\sigma(F^2)] = 0.040$

$wR(F^2) = 0.112$

$S = 1.07$

1654 reflections

128 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-atom parameters constrained

$w = 1/[\sigma^2(F_o^2) + (0.0644P)^2 + 0.0983P]$

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

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

$\Delta\rho_{\max} = 0.20$ e Å⁻³

$\Delta\rho_{\min} = -0.34$ 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	$U_{\text{iso}}^*/U_{\text{eq}}$
O1	1.00611 (13)	1.92652 (14)	1.27797 (14)	0.0228 (3)
O2	0.62238 (13)	1.01503 (13)	0.83520 (13)	0.0192 (3)
C1	1.22612 (19)	1.8825 (2)	1.4939 (2)	0.0246 (3)
H1A	1.2907	2.0214	1.5498	0.037*
H1B	1.3039	1.8086	1.4524	0.037*
H1C	1.1976	1.8627	1.5904	0.037*
C2	1.05129 (18)	1.8131 (2)	1.32364 (19)	0.0179 (3)
C3	0.93621 (18)	1.60305 (19)	1.20855 (18)	0.0166 (3)
C4	0.97991 (18)	1.4674 (2)	1.25479 (18)	0.0174 (3)
H4	1.0826	1.5113	1.3695	0.021*
C5	0.87715 (18)	1.2701 (2)	1.13753 (19)	0.0174 (3)
H5	0.9082	1.1802	1.1722	0.021*
C6	0.72738 (18)	1.20521 (19)	0.96773 (18)	0.0166 (3)
C7	0.67704 (18)	1.3397 (2)	0.92338 (19)	0.0181 (3)
H7	0.5714	1.2964	0.8115	0.022*
C8	0.78030 (19)	1.5347 (2)	1.04150 (19)	0.0181 (3)
H8	0.7455	1.6249	1.0095	0.022*
C9	0.67885 (19)	0.86901 (19)	0.86356 (19)	0.0197 (3)
H9A	0.6449	0.8585	0.9629	0.024*
H9B	0.8132	0.9049	0.9071	0.024*
C10	0.58484 (18)	0.67807 (19)	0.67576 (19)	0.0172 (3)
C11	0.64548 (18)	0.64417 (19)	0.52994 (19)	0.0183 (3)
H11	0.7453	0.7424	0.5498	0.022*
C12	0.56143 (18)	0.4684 (2)	0.35604 (19)	0.0176 (3)
H12	0.6038	0.4474	0.2576	0.021*

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
O1	0.0258 (6)	0.0168 (5)	0.0241 (5)	0.0062 (4)	0.0077 (4)	0.0111 (4)
O2	0.0205 (5)	0.0115 (5)	0.0182 (5)	0.0045 (4)	0.0024 (4)	0.0057 (4)
C1	0.0232 (7)	0.0162 (7)	0.0236 (7)	0.0013 (6)	0.0030 (6)	0.0080 (6)
C2	0.0192 (7)	0.0161 (7)	0.0177 (7)	0.0052 (6)	0.0089 (6)	0.0079 (6)
C3	0.0171 (7)	0.0161 (7)	0.0164 (7)	0.0052 (6)	0.0080 (5)	0.0078 (6)
C4	0.0174 (7)	0.0185 (7)	0.0137 (6)	0.0064 (5)	0.0050 (5)	0.0071 (6)
C5	0.0206 (7)	0.0160 (7)	0.0178 (7)	0.0079 (6)	0.0078 (6)	0.0099 (6)
C6	0.0178 (7)	0.0143 (7)	0.0158 (7)	0.0041 (5)	0.0076 (6)	0.0064 (6)
C7	0.0181 (7)	0.0179 (7)	0.0151 (6)	0.0056 (6)	0.0041 (5)	0.0078 (6)
C8	0.0207 (7)	0.0172 (7)	0.0190 (7)	0.0080 (6)	0.0078 (6)	0.0111 (6)
C9	0.0226 (7)	0.0154 (7)	0.0190 (7)	0.0077 (6)	0.0046 (6)	0.0091 (6)

C10	0.0184 (7)	0.0143 (7)	0.0190 (7)	0.0080 (5)	0.0050 (5)	0.0094 (6)
C11	0.0168 (7)	0.0165 (7)	0.0225 (7)	0.0050 (5)	0.0060 (5)	0.0124 (6)
C12	0.0194 (7)	0.0184 (7)	0.0189 (7)	0.0089 (6)	0.0083 (5)	0.0115 (6)

Geometric parameters (Å, °)

O1—C2	1.2240 (16)	C6—C7	1.3967 (19)
O2—C6	1.3617 (17)	C7—C8	1.375 (2)
O2—C9	1.4398 (15)	C7—H7	0.9500
C1—C2	1.5059 (19)	C8—H8	0.9500
C1—H1A	0.9800	C9—C10	1.5020 (18)
C1—H1B	0.9800	C9—H9A	0.9900
C1—H1C	0.9800	C9—H9B	0.9900
C2—C3	1.4866 (19)	C10—C11	1.3931 (19)
C3—C4	1.3964 (19)	C10—C12 ⁱ	1.3954 (19)
C3—C8	1.4035 (19)	C11—C12	1.3864 (19)
C4—C5	1.387 (2)	C11—H11	0.9500
C4—H4	0.9500	C12—C10 ⁱ	1.3954 (19)
C5—C6	1.3972 (19)	C12—H12	0.9500
C5—H5	0.9500		
C6—O2—C9	117.64 (10)	C8—C7—C6	120.06 (12)
C2—C1—H1A	109.5	C8—C7—H7	120.0
C2—C1—H1B	109.5	C6—C7—H7	120.0
H1A—C1—H1B	109.5	C7—C8—C3	121.19 (13)
C2—C1—H1C	109.5	C7—C8—H8	119.4
H1A—C1—H1C	109.5	C3—C8—H8	119.4
H1B—C1—H1C	109.5	O2—C9—C10	108.13 (10)
O1—C2—C3	120.37 (12)	O2—C9—H9A	110.1
O1—C2—C1	120.78 (12)	C10—C9—H9A	110.1
C3—C2—C1	118.84 (12)	O2—C9—H9B	110.1
C4—C3—C8	117.90 (13)	C10—C9—H9B	110.1
C4—C3—C2	122.90 (12)	H9A—C9—H9B	108.4
C8—C3—C2	119.15 (12)	C11—C10—C12 ⁱ	118.81 (13)
C5—C4—C3	121.68 (12)	C11—C10—C9	119.68 (13)
C5—C4—H4	119.2	C12 ⁱ —C10—C9	121.51 (12)
C3—C4—H4	119.2	C12—C11—C10	120.58 (13)
C4—C5—C6	119.14 (13)	C12—C11—H11	119.7
C4—C5—H5	120.4	C10—C11—H11	119.7
C6—C5—H5	120.4	C11—C12—C10 ⁱ	120.60 (12)
O2—C6—C7	115.22 (12)	C11—C12—H12	119.7
O2—C6—C5	124.87 (12)	C10 ⁱ —C12—H12	119.7
C7—C6—C5	119.91 (13)		

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

Hydrogen-bond geometry (Å, °)

Cg1 is the centroid of the C3–C8 ring.

<i>D</i> —H··· <i>A</i>	<i>D</i> —H	H··· <i>A</i>	<i>D</i> ··· <i>A</i>	<i>D</i> —H··· <i>A</i>
C11—H11···O1 ⁱⁱ	0.95	2.54	3.4362 (18)	158
C12—H12···Cg1 ⁱⁱⁱ	0.95	2.61	3.5078 (17)	158

Symmetry codes: (ii) $-x+2, -y+3, -z+2$; (iii) $x, y-1, z-1$.