## organic compounds

Acta Crystallographica Section E

**Structure Reports** 

**Online** 

ISSN 1600-5368

## (2-Methylphenyl)(phenyl)methanol

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Received 21 July 2010; accepted 23 July 2010

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Key indicators: single-crystal X-ray study; T = 100 K; mean  $\sigma(C-C) = 0.002$  Å; R factor = 0.045; wR factor = 0.121; data-to-parameter ratio = 17.6.

In the title compound,  $C_{14}H_{14}O$ , the two benzene rings are almost orthogonal [dihedral angle = 87.78 (8) °]. The hydroxy group lies approximately in the plane of its attached benzene ring [O-C-C-C torsion angle =  $-17.47 (17)^{\circ}$ ], and the hydroxyl and methyl groups lie to the same side of the molecule and are *gauche* to each other. In the crystal, a hexameric aggregate mediated by a ring of six  $O-H\cdots O$  hydrogen bonds occurs, generating an  $R_6^6(12)$  loop.

#### **Related literature**

For general background to the use of benzhydrols, see: Ohkuma *et al.* (2000). For the use of the title compound in the perfume and pharmaceutical industries, see: Meguro *et al.* (1985). For related diphenylmethanol structures, see: Ferguson *et al.* (1995).

#### **Experimental**

Crystal data

 $C_{14}H_{14}O$   $M_r = 198.25$ Trigonal,  $R\overline{3}$  a = 23.013 (2) Å c = 10.6067 (11) Å $V = 4864.8 (7) \text{ Å}^3$  Z = 18Mo  $K\alpha$  radiation  $\mu = 0.08 \text{ mm}^{-1}$  T = 100 K $0.40 \times 0.35 \times 0.30 \text{ mm}$ 

Data collection

Bruker SMART APEX CCD diffractometer Absorption correction: multi-scan (SADABS; Sheldrick, 1996)  $T_{\min} = 0.971, T_{\max} = 0.978$ 

6286 measured reflections 2475 independent reflections 2022 reflections with  $I > 2\sigma(I)$   $R_{\rm int} = 0.026$ 

Refinement

 $R[F^2 > 2\sigma(F^2)] = 0.045$   $wR(F^2) = 0.121$  S = 1.082475 reflections 141 parameters 1 restraint

H atoms treated by a mixture of independent and constrained refinement

 $\Delta \rho_{\text{max}} = 0.43 \text{ e Å}^{-3}$  $\Delta \rho_{\text{min}} = -0.30 \text{ e Å}^{-3}$ 

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

$D$ $ H$ $\cdot \cdot \cdot A$	D-H	$H \cdot \cdot \cdot A$	$D \cdot \cdot \cdot A$	$D-\mathrm{H}\cdots A$
O1-H1···O1 <sup>i</sup>	0.85 (1)	1.85 (1)	2.6967 (10)	174 (2)

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

Data collection: *APEX2* (Bruker, 2008); cell refinement: *SAINT* (Bruker, 2008); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *ORTEP-3* (Farrugia, 1997) and *DIAMOND* (Brandenburg, 2006); software used to prepare material for publication: *publCIF* (Westrip, 2010).

HSY thanks the University of Mysore for research facilities and for sabbatical leave. BPS thanks R. L. Fine Chemicals for the gift of the title compound. The authors are also grateful to the University of Malaya for support of the crystallographic facility.

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: HB5570).

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supplementary m	aterials	

Acta Cryst. (2010). E66, o2136 [doi:10.1107/S1600536810029417]

## (2-Methylphenyl)(phenyl)methanol

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#### Comment

Benzhydrols are widely used as intermediates for the synthesis of pharmaceuticals (Ohkuma *et al.*, 2000), including drugs such as diphenhydramine, orphenadrine, diphenidol and phenyltoloxamine. The crystal structures and hydrogen bonding in some diphenylmethanols have been reported (Ferguson *et al.*, 1995). The title compound, phenyl-*o*-tolyl-methanol, (I), is a derivative of diphenylmethanol and it has use in the perfume and pharmaceutical industries (Meguro *et al.*, 1985).

The molecular structure of (I), Fig. 1, features a tertiary C7 atom connected to benzene and o-tolyl rings. With reference to the benzene ring, the O1 atom is nearly co-planar as seen in the O1–C8–C9–C14 torsion angle of -17.47 (17) °. By contrast, the o-tolyl group is almost orthogonal as seen in the C1–C8–C9–C10 torsion angle of -80.12 (15) °; the dihedral angle formed between the two least-squares planes is 87.78 (8) °. While lying to the same side of the molecule, the hydroxy and methyl groups are *gauche*.

The crystal packing is dominated by O–H···O hydrogen bonding, Table 1. Almost planar 12-membered rings comprising six O–H···O hydrogen bonds are found, each disposed about a site of symmetry,  $\overline{3}$ , Fig. 2. The hexameric aggregates stack in columns aligned along the c axis, Fig. 3.

#### **Experimental**

The title compound was obtained as a gift from R. L. Fine Chemicals, Bangalore, India. Colourless blocks of (I) were obtained by the slow evaporation of its acetonitrile solution; m.pt. 369–372 K.

#### Refinement

Carbon-bound H-atoms were placed in calculated positions (C—H 0.95 to 1.00 Å) and were included in the refinement in the riding model approximation, with  $U_{iso}(H)$  set to 1.2 to 1.5 $U_{equiv}(C)$ . The O-bound H-atom was located in a difference Fourier map, and was refined with a distance restraint of O–H 0.84±0.01 Å; the  $U_{iso}$  value was freely refined

### **Figures**

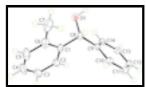


Fig. 1. The molecular structure of (I) showing displacement ellipsoids at the 50% probability level.

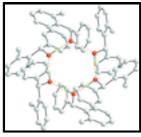


Fig. 2. A hexameric aggregate mediated by O–H···O hydrogen bonds (orange dashed lines) in (I). Non-participating H atoms have been omitted.



Fig. 3. The unit-cell contents shown in projection down the c axis in (I). The O–H···O hydrogen bonding is shown as orange dashed lines.

## (2-Methylphenyl)(phenyl)methanol

Crystal data

 $C_{14}H_{14}O$   $D_x = 1.218 \text{ Mg m}^{-3}$ 

 $M_r = 198.25$  Mo  $K\alpha$  radiation,  $\lambda = 0.71073$  Å Trigonal,  $R\overline{3}$  Cell parameters from 2551 reflections

Hall symbol: -R 3  $\theta = 2.8-28.3^{\circ}$  a = 23.013 (2) Å  $\mu = 0.08 \text{ mm}^{-1}$ 

c = 10.6067 (11) Å T = 100 K $V = 4864.8 (7) \text{ Å}^3$  Block, colourless

Z = 18  $0.40 \times 0.35 \times 0.30 \text{ mm}$  F(000) = 1908

Data collection

Bruker SMART APEX CCD 2475 independent reflections

diffractometer 2473 independent reflections Radiation source: fine-focus sealed tube 2022 reflections with  $I > 2\sigma(I)$ 

graphite  $R_{\text{int}} = 0.026$ 

θ<sub>max</sub> = 27.5°, θ<sub>min</sub> = 1.8°Absorption correction: multi-scan

Absorption correction: multi-scall  $h = -28 \rightarrow 28$  (SADABS; Sheldrick, 1996)  $T_{\text{min}} = 0.971$ ,  $T_{\text{max}} = 0.978$   $k = -29 \rightarrow 14$  6286 measured reflections  $l = -13 \rightarrow 13$ 

Refinement

 $R[F^2 > 2\sigma(F^2)] = 0.045$ 

Refinement on  $F^2$  Primary atom site location: structure-invariant direct

methods

Least-squares matrix: full Secondary atom site location: difference Fourier map

Hydrogen site location: inferred from neighbouring

sites

$wR(F^2) = 0.121$	H atoms treated by a mixture of independent and constrained refinement
S = 1.08	$w = 1/[\sigma^2(F_0^2) + (0.0533P)^2 + 4.2609P]$ where $P = (F_0^2 + 2F_c^2)/3$
2475 reflections	$(\Delta/\sigma)_{\text{max}} < 0.001$
141 parameters	$\Delta \rho_{max} = 0.43 \text{ e Å}^{-3}$
1 restraint	$\Delta \rho_{min} = -0.30 \text{ e Å}^{-3}$

### 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  $(\mathring{A}^2)$ 

	x	У	z	$U_{\rm iso}*/U_{\rm eq}$
O1	0.13170 (5)	0.08915 (5)	0.01453 (9)	0.0229(2)
H1	0.1201 (9)	0.0483 (5)	0.0010 (18)	0.043 (5)*
C1	0.20865 (7)	0.19098 (7)	0.11187 (14)	0.0237(3)
C2	0.18864 (8)	0.21209 (7)	0.21715 (15)	0.0306(3)
H2	0.1637	0.1808	0.2816	0.037*
C3	0.20471 (9)	0.27864 (8)	0.22928 (18)	0.0417 (4)
Н3	0.1911	0.2930	0.3018	0.050*
C4	0.24081 (9)	0.32372 (8)	0.1346 (2)	0.0495 (5)
H4	0.2523	0.3694	0.1423	0.059*
C5	0.26018 (8)	0.30288 (8)	0.0295 (2)	0.0433 (5)
H5	0.2847	0.3344	-0.0349	0.052*
C6	0.24463 (7)	0.23669 (7)	0.01529 (16)	0.0321 (4)
C7	0.26458 (8)	0.21578 (9)	-0.10289 (17)	0.0422 (4)
H7A	0.2878	0.2545	-0.1591	0.063*
H7B	0.2244	0.1806	-0.1450	0.063*
H7C	0.2946	0.1984	-0.0819	0.063*
C8	0.18897 (6)	0.11780 (6)	0.09612 (12)	0.0206(3)
H8	0.2264	0.1162	0.0515	0.025*
C9	0.17524 (6)	0.07865 (6)	0.21775 (12)	0.0183 (3)
C10	0.22851 (7)	0.08429 (7)	0.28925 (13)	0.0247 (3)
H10	0.2733	0.1148	0.2642	0.030*
C11	0.21660 (7)	0.04561 (7)	0.39708 (14)	0.0276(3)
H11	0.2533	0.0500	0.4457	0.033*
C12	0.15144 (7)	0.00071 (7)	0.43395 (13)	0.0247 (3)
H12	0.1433	-0.0263	0.5069	0.030*

C13 H13 C14 H14	0.09838 (7) 0.0536 0.11032 (7) 0.0735	-0.00461 (7 -0.0350 0.03426 (6) 0.0304	0.3894	32 (13)	0.0229 (3) 0.027* 0.0207 (3) 0.025*		
Atomic displa	acement parameters	$(\mathring{A}^2)$					
	$U^{11}$	$U^{22}$	$U^{33}$	$U^{12}$	$U^{13}$	$U^{23}$	
O1	0.0260 (5)	0.0200 (5)	0.0247 (5)	0.0130(4)	-0.0060(4)	-0.0017 (4)	
C1	0.0180 (6)	0.0186 (6)	0.0340 (8)	0.0087 (5)	-0.0060 (5)	0.0017 (5)	
C2	0.0317 (8)	0.0235 (7)	0.0397 (9)	0.0161 (6)	-0.0096 (6)	-0.0041 (6)	
C3	0.0449 (10)	0.0297 (8)	0.0577 (11)	0.0241 (8)	-0.0221 (8)	-0.0149 (8)	
C4	0.0386 (10)	0.0164 (7)	0.0895 (15)	0.0106 (7)	-0.0297 (10)	-0.0038 (8)	
C5	0.0261 (8)	0.0226 (8)	0.0717 (13)	0.0051 (6)	-0.0108 (8)	0.0127 (8)	
C6	0.0162 (6)	0.0257 (7)	0.0487 (10)	0.0062 (6)	-0.0050(6)	0.0106 (7)	
C7	0.0275 (8)	0.0477 (10)	0.0491 (10)	0.0171 (8)	0.0090(7)	0.0230(8)	
C8	0.0190(6)	0.0213 (6)	0.0239 (7)	0.0118 (5)	-0.0006(5)	0.0022 (5)	
C9	0.0205 (6)	0.0158 (6)	0.0209 (6)	0.0107 (5)	-0.0001 (5)	-0.0007 (5)	
C10	0.0188 (6)	0.0233 (7)	0.0301 (7)	0.0091 (5)	-0.0017(5)	0.0023 (5)	
C11	0.0246 (7)	0.0302 (8)	0.0289 (8)	0.0144 (6)	-0.0055 (6)	0.0027 (6)	
C12	0.0301 (7)	0.0228 (7)	0.0234 (7)	0.0148 (6)	-0.0001 (5)	0.0027 (5)	
C13	0.0220 (7)	0.0217 (6)	0.0237 (7)	0.0100 (5)	0.0020 (5)	-0.0008(5)	
C14	0.0193 (6)	0.0218 (6)	0.0230 (7)	0.0118 (5)	-0.0014 (5)	-0.0016 (5)	
Geometric na	vrameters (Å, °)						
		1 4222 (16)	67 1	17D	0.00	00	
O1—C8		1.4323 (16) 0.852 (9)	С7—I С7—I		0.98 0.98		
O1—H1 C1—C2			C7—r				
C1—C2 C1—C6		1.385 (2) 1.404 (2)			1.31	37 (18)	
C1—C0 C1—C8		1.5188 (18)				62 (18)	
C1—C6 C2—C3		1.390 (2)					
C2—H2		0.9500				1.3911 (18) 1.390 (2)	
C3—C4		1.383 (3)				0.9500	
C3—H3		0.9500	C10—H10 C11—C12		1.386 (2)		
C4—C5		1.373 (3)	C11—		0.9500		
C4—H4		0.9500	C12—C13		1.3806 (19)		
C5—C6		1.388 (2)	C12—		0.95		
C5—H5		0.9500	C13—		1.3868 (19)		
C6—C7		1.495 (3)	C13—H13		0.9500		
C7—H7A		0.9800	C14—	-H14	0.95	00	
C8—O1—H1		108.5 (13)	01—0	C8—C9	111.	77 (10)	
C2—C1—C6		120.03 (13)		C8—C1		81 (10)	
C2—C1—C8		120.69 (13)		C8—C1		10 (11)	
C6—C1—C8		119.20 (13)		C8—H8	108.		
C1—C2—C3		120.59 (16)		C8—H8	108.		
C1—C2—H2		119.7	C1—C	C8—H8	108.	0	
C3—C2—H2		119.7	C14—	-C9—C10	118.	73 (12)	

C4 C2 C2	110 22 (10)	C14 C0 C0		121 27 (12)
C4—C3—C2 C4—C3—H3	119.23 (18) 120.4	C14—C9—C8 C10—C9—C8		121.37 (12) 119.81 (11)
C2—C3—H3	120.4	C10—C9—C8 C11—C10—C9		120.43 (13)
				` ′
C5—C4—C3	120.38 (15)	C11—C10—H10		119.8
C5—C4—H4	119.8	C9—C10—H10		119.8
C3—C4—H4	119.8	C12—C11—C10		120.20 (13)
C4—C5—C6	121.39 (17)	C12—C11—H11		119.9
C4—C5—H5	119.3	C10—C11—H11		119.9
C6—C5—H5	119.3	C13—C12—C11		119.60 (13)
C5—C6—C1	118.37 (16)	C13—C12—H12		120.2
C5—C6—C7	119.48 (15)	C11—C12—H12		120.2
C1—C6—C7	122.12 (14)	C12—C13—C14		120.14 (13)
C6—C7—H7A	109.5	C12—C13—H13		119.9
C6—C7—H7B	109.5	C14—C13—H13		119.9
H7A—C7—H7B	109.5	C9—C14—C13		120.89 (12)
C6—C7—H7C	109.5	C9—C14—H14		119.6
H7A—C7—H7C	109.5	C13—C14—H14		119.6
H7B—C7—H7C	109.5			
C6—C1—C2—C3	-1.0 (2)	C6—C1—C8—C9		157.52 (12)
C8—C1—C2—C3	-177.73 (13)	O1—C8—C9—C14		-17.47 (17)
C1—C2—C3—C4	0.3 (2)	C1—C8—C9—C14		103.27 (14)
C2—C3—C4—C5	0.4(2)	O1—C8—C9—C10		159.14 (12)
C3—C4—C5—C6	-0.3 (2)	C1—C8—C9—C10		-80.12 (15)
C4—C5—C6—C1	-0.4(2)	C14—C9—C10—C11		0.6(2)
C4—C5—C6—C7	177.76 (15)	C8-C9-C10-C11		-176.11 (12)
C2—C1—C6—C5	1.0(2)	C9—C10—C11—C12		0.4(2)
C8—C1—C6—C5	177.84 (13)	C10—C11—C12—C13		-1.0(2)
C2—C1—C6—C7	-177.07 (14)	C11—C12—C13—C14		0.7(2)
C8—C1—C6—C7	-0.3 (2)	C10—C9—C14—C13		-0.92 (19)
C2—C1—C8—O1	98.25 (14)	C8—C9—C14—C13		175.72 (12)
C6—C1—C8—O1	-78.54 (14)	C12—C13—C14—C9		0.3 (2)
C2—C1—C8—C9	-25.70 (17)			
Hydrogen-bond geometry (Å, °)				
<i>D</i> —H··· <i>A</i>	<i>D</i> —H	$H\cdots A$	D··· $A$	D— $H$ ··· $A$

0.85(1)

1.85 (1)

2.6967 (10)

174.(2)

O1—H1···O1<sup>i</sup>

Symmetry codes: (i) y, -x+y, -z.

Fig. 1

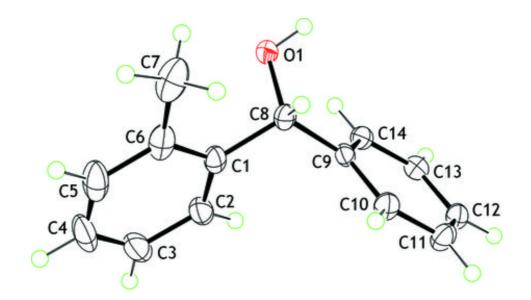


Fig. 2

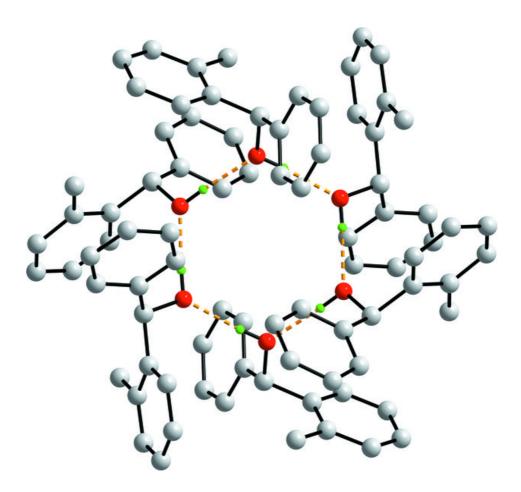


Fig. 3

