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#### **Key indicators**

Single-crystal X-ray study T = 292 KMean  $\sigma(\text{C}-\text{C}) = 0.002 \text{ Å}$  R factor = 0.045 wR factor = 0.120 Data-to-parameter ratio = 17.7

For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e.

## 1,2-Dimethyl-4,5-bis(phenoxymethyl)benzene

The molecule of the title compound,  $C_{22}H_{22}O_2$ , possesses a crystallographically imposed twofold axis. The two terminal phenyl rings both make a dihedral angle of 82.5 (2)° with the central benzene ring. The crystal packing is stabilized mainly by van der Waals forces.

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#### Comment

The title compound, (I), is a precursor to substituted phenylene-vinylene oligomers, which are novel compounds for use as semiconductors in an electronic nose (Vanneste *et al.*, 1998).



The molecule of (I) possesses a crystallographically imposed twofold axis (Fig. 1). The bond lengths and angles are within normal ranges (Table 1) (Allen *et al.*, 1987). The torsion angles C3-C4-C5-O1 and C4-C5-O1-C6 are 90.40 (15) and 178.23 (11)°, respectively. The two terminal phenyl rings both make dihedral angles of 82.5 (2)° with the central benzene ring, while the dihedral angle between the phenyl rings is 56.8 (1)°.

The crystal packing (Fig. 2) is stabilized mainly by van der Waals forces.

### **Experimental**

Sodium hydride (0.48 g, 20 mmol) was added to a stirred solution of 1,2-bis-chloromethyl-4,5-dimethylbenzene (2.03 g, 10 mmol) in anhydrous tetrahydrofuran (THF, 30 ml) under a nitrogen atmosphere. To this mixture, a solution of phenol (1.88 g, 20 mmol) in anhydrous THF (10 ml) was added dropwise and stirred overnight at room temperature. The THF was removed and water was added to it, followed by extraction with dichloromethane. The solvent was evaporated to dryness and the compound was purified by column chromatography to obtain the title compound (yield 2.86 g, 90%). A sample was dissolved in a mixture of hexane–EtOAc (1:6) at room temperature and ambient pressure, and crystals of (I) grew over a period of a week when the solution was exposed to the air.

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#### Figure 1

A view of (I), showing the atom-labelling scheme [symmetry code: (a)  $1 - x, y, \frac{1}{2} - z$ ]. Displacement ellipsoids are drawn at the 50% probability level.



#### Figure 2

The crystal packing, viewed approximately along the b axis.

Crystal data

C22H22O2 Mo  $K\alpha$  radiation  $M_r = 318.40$ Cell parameters from 1978 Orthorhombic, Pbcn reflections a = 18.476 (4) Å  $\theta=2.4{-}21.6^\circ$  $\mu = 0.07~\mathrm{mm}^{-1}$ b = 11.494 (2) Å c = 8.4632 (17) Å T = 292 (2) K V = 1797.2 (6) Å<sup>3</sup> Block, colourless Z = 4 $0.25 \times 0.20 \times 0.20$  mm  $D_x = 1.177 \text{ Mg m}^{-3}$ 

Data collection

Bruker SMART CCD area-detector	1180 reflections with $I > 2\sigma(I)$	
diffractometer	$R_{\rm int} = 0.057$	
$\varphi$ and $\omega$ scans	$\theta_{\rm max} = 27.0^{\circ}$	
Absorption correction: none	$h = -15 \rightarrow 23$	
11066 measured reflections	$k = -14 \rightarrow 14$	
1964 independent reflections	$l = -10 \rightarrow 10$	
Refinement		
Refinement on $F^2$	$w = 1/[\sigma^2(F_0^2) + (0.065P)^2]$	
$R[F^2 > 2\sigma(F^2)] = 0.045$	where $P = (F_0^2 + 2F_c^2)/3$	
$wR(F^2) = 0.120$	$(\Delta/\sigma)_{\rm max} < 0.001$	
S = 0.89	$\Delta \rho_{\rm max} = 0.16 \ {\rm e} \ {\rm \AA}^{-3}$	
1964 reflections	$\Delta \rho_{\rm min} = -0.14 \text{ e} \text{ Å}^{-3}$	
111 parameters	Extinction correction: SHELX97	
H-atom parameters constrained	(Sheldrick, 1997)	
	Extinction coefficient: 0.0048 (13)	

Table 1Selected geometric parameters (Å, °).

C4-C5	1.5011 (18)	C6-O1	1.3746 (16)
C5-O1	1.4329 (16)		
C3-C4-C5	119.76 (13)	O1-C6-C7	115.68 (13)
O1-C5-C4	108.42 (11)	C6-O1-C5	116.84 (10)

After their location in a difference Fourier map, H atoms were placed in calculated positions and allowed to ride on their parent atoms, with C-H = 0.93-0.97 Å and  $U_{\rm iso}(\rm H) = 1.2-1.5~U_{eq}(\rm C)$ .

Data collection: *SMART* (Bruker, 1997); cell refinement: *SMART*; data reduction: *SAINT* (Bruker, 1999); program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELX97* (Sheldrick, 1997); molecular graphics: *SHELXTL* (Bruker, 2001); software used to prepare material for publication: *SHELXTL*.

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