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#### **Key indicators**

Single-crystal X-ray study T = 293 KMean  $\sigma(C-C) = 0.007 \text{ Å}$  R factor = 0.029 wR factor = 0.056 Data-to-parameter ratio = 17.4

For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e. 2,4-Dibromophenyl 2,6-dibromophenyl ether

The title compound,  $C_{12}H_6Br_4O$ , is the fourth well characterized of a total of potentially 209 different brominated diphenyl ethers. Salient intermolecular interactions are between ether O and aromatic C atoms, and between a bromine substituent and the aromatic ring system.

#### Comment

The title compound, (I), belongs to a class of compounds known as brominated diphenyl ethers (DE's) which are used as flame retardants. Commercially available mixtures mainly consist of highly brominated congeners, such as decabromodiphenyl ether (Eriksson *et al.*, 1999). The occurrence in the environment of less brominated congeners is dependent on both primary sources and decomposition from higher brominated DE's (Eriksson *et al.*, 2001). The crystal structures and packing patterns of these compounds are of fundamental importance in order to model the reaction mechanism in solidstate reaction, *e.g.* flame retardants adsorbed on soot particles.



Both aromatic rings of the title compound are planar within less than 0.01 Å. The deviation of the substituents from the ring plane of the first ring (C1–C6) are 0.050 (7) Å for O and 0.031 (8) Å for Br4, while Br3 is within the ring plane. The substituents of the second ring (C7–C12) all deviate significantly from the ring plane: O 0.016 (6) Å, Br1 0.110 (7) Å and Br2 0.053 (7) Å. The angle between the two ring planes is  $89.1 (2)^{\circ}$ .

Designating the aromatic rings of the molecule as ring1 = C1-C6 and ring2 = C7-C12, the structure can be described as a packing of molecules with close contacts between symmetry-equivalent ring1 systems (C1-C6)···(C1-C6)<sup>i</sup> etc. [symmetry code: (i) -0.5-x, 0.5+y, -0.5-z]. Each of the neighbouring symmetry-equivalent ring1's of the molecules are inclined at 37.9 (1)° to each other (Fig. 2). The two shortest intermolecular C···O contacts are O-C3<sup>ii</sup> and O···C4<sup>ii</sup>, both of these distances being 3.33 (1) Å [symmetry code: (ii) -x-0.5, y-0.5, -z+0.5]. These intermolecular C···O distances are quite short compared to previously known structures containing the diphenyl ether moiety from the Cambridge

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### Figure 1

One molecule of the title compound, with the atom-numbering scheme. Displacement ellipsoids are shown at the 50% probability level. H atoms are shown as small circles of arbitrary radii.

Structural Database (CSD; Allen & Kennard, 1993). Another short intermolecular contact is due to ring2 packing with one parallel symmetry-equivalent copy of itself on one side together with a short  $Br \cdot \cdot C$  interaction on the opposite side of ring2 [Br4...C11<sup>iii</sup> = 3.548 (5) Å; symmetry code: (iii) x-1, y, z]; this is the shortest intermolecular  $Br \cdots C$  contact distance in the present structure.

Thus, the structure can be described as a packing of long chains held together by intermolecular interactions between ring1 of different molecules; ring2 then links different chains to each other. Intermolecular Br...Br contacts further stabilize the structure, with the shortest intermolecular  $Br \cdots Br$ distance being  $Br3 \cdot \cdot \cdot Br4^{iv} = 3.798$  (1) Å [symmetry code: (iv) x+1, y, z]. This is not a specially short intermolecular distance compared to other bromine substituted aromatic compounds found in the CSD and is significantly longer than the short Br...Br contact in phenyl 2,4,6-tribromophenyl ether (Eriksson & Hu, 2001). Thus, one can speculate that the  $Br \cdot \cdot \cdot Br$  contacts only contribute a small part of the lattice energy.

## **Experimental**

The synthesis of the PBDE was carried out by coupling a diphenyliodonium salt with a bromophenylate (Beringer et al., 1959; Ziegler & Marr, 1962; Hu, 1996, 1999). The title compound was recrystallized from methanol.

### Crystal data

$C_{12}H_6Br_4O$	$D_{\rm r} = 2.331 {\rm Mg m}^{-3}$
$M_r = 485.81$	Mo $K\alpha$ radiation
Monoclinic, $P2_1/n$	Cell parameters from 1520
$a = 8.0274 (12) \text{\AA}$	reflections
b = 8.389 (2) Å	$\theta = 2.0-26.0^{\circ}$
c = 20.562 (3)  Å	$\mu = 11.61 \text{ mm}^{-1}$
$\beta = 90.606 \ (19)^{\circ}$	T = 293 (2) K
$V = 1384.5(5) \text{ Å}^3$	Prism, colourless
Z = 4	$0.19 \times 0.13 \times 0.10 \text{ mm}$



#### Figure 2

Stereoview of the molecular packing, with the close contacts between one of the two different aromatic rings (C1-C6) and symmetry equivalents.

Data collection	
Stoe IPDS diffractometer	1625 reflections with $I > 2\sigma(I)$
$\varphi$ scans	$R_{\rm int} = 0.049$
Absorption correction: numerical	$\theta_{\rm max} = 26.0^{\circ}$
(X-RED; Stoe & Cie, 1998)	$h = -9 \rightarrow 9$
$T_{\min} = 0.113, \ T_{\max} = 0.307$	$k = -10 \rightarrow 10$
21426 measured reflections	$l = -25 \rightarrow 25$
2676 independent reflections	
Refinement	

Refinement on $F^2$	H-atom parameters constrained
$R[F^2 > 2\sigma(F^2)] = 0.029$	$w = 1/[\sigma^2(F_o^2) + (0.01P)^2]$
$wR(F^2) = 0.056$	where $P = (F_o^2 + 2F_c^2)/3$
S = 0.98	$(\Delta/\sigma)_{\rm max} = 0.001$
2676 reflections	$\Delta \rho_{\rm max} = 0.73 \ {\rm e} \ {\rm \AA}^{-3}$
154 parameters	$\Delta \rho_{\rm min} = -0.48 \text{ e } \text{\AA}^{-3}$

# Table 1

Selected geometric parameters (Å, °).

Br1-C10	1.895 (5)	Br4-C2	1.879 (5)
Br2-C12	1.902 (5)	O-C7	1.395 (5)
Br3-C6	1.885 (5)	O-C1	1.397 (5)
C7-O-C1	115.7 (4)	C8-C7-O	123.3 (4)
C6-C1-O	122.2 (4)	C12-C7-O	116.9 (5)
O-C1-C2	118.8 (4)		
C1-O-C7-C8	26.3 (6)	C1-O-C7-C12	-154.2 (4)
C7-O-C1-C2	-104.6(5)	C7-O-C1-C6	78.0 (6)

Two data sets were collected with the Stoe IPDS system (Stoe & Cie, 1997), and these were merged and scaled together. The applied scale factor between the two data sets was 0.989.

Data collection: EXPOSE (Stoe & Cie, 1997); cell refinement: CELL (Stoe & Cie, 1997); data reduction: INTEGRATE (Stoe & Cie, 1997) and X-RED (Stoe & Cie, 1997); program(s) used to solve structure: SHELXS97 (Sheldrick, 1990); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: DIAMOND (Bergerhoff, 1996).

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