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Key indicators

Single-crystal X-ray study T = 293 K Mean σ (C–C) = 0.010 Å R factor = 0.036 wR factor = 0.066 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. The title compound, $C_{12}H_5Br_5O$, belongs to a group of flame retardants known as polybrominated diphenyl ethers (PBDE). Salient features of the packing are stabilization due to intermolecular $Br \cdots Br$ contacts in the *ab* plane and aromatic intermolecular contacts along the *c* direction.

2,3,4,5,6-Pentabromophenyl phenyl ether

Comment

An important group of flame retardants are the polybrominated diphenyl ethers (PBDE). There are a total of 209 different PBDE's. The modelling of the reactivity of different PBDE's is a task that requires accurate geometries of the molecular species. Geometric data from crystallographic measurements on brominated diphenyl ethers without hetero substituents other than bromine are rather limited (Örn *et al.*, 1996; Eriksson *et al.*, 1999; Mrse *et al.*, 2000; Eriksson & Hu, 2001, 2002; Eriksson *et al.*, 2002*a,b*). In addition, a partial structure of bis(4-bromophenyl) ether (Toussaint, 1946) has been published. In a search of the Cambridge Structural Database (Allen & Kennard, 1993), including hetero substituents other than bromine, a larger set of structures for use as model compounds was found, but still only in the order of 10– 15 different structures.



The bromine-substituted ring is planar with an r.m.s. deviation of 0.012 Å for the C atoms (C1–C6) of the ring. The O atom deviates most from this plane, by 0.163 (8) Å, while atoms Br1, Br2 and Br3 lie within 2σ of the plane; atoms Br4 and Br5 deviate by 0.037 (8) and 0.035 (8) Å, respectively. The second, unsubstituted, aromatic ring (C7–C12) is more strictly planar, with an r.m.s. deviation of 0.004 Å for the C atoms. The O atom is coplanar with this ring. The angle between the two ring planes is 89.7 (2)°.

A view of the arrangement of the molecules of the title compound, (I), is shown in Fig. 2, where the two shortest intermolecular Br···Br contacts are marked. The closest intermolecular Br···Br contacts are: Br3···Br4ⁱ = 3.567 (1) Å and Br2···Br2ⁱⁱ = 3.698 (1) Å [symmetry codes: (i) -x, -1/2+y, 1/2-z; (ii) 1-x, 1/2+y, 1/2-z]. The Br2 atoms of different molecules form a connected zigzag chain of close contacts along the *b* axis. Similarly, atoms Br3 and Br4 form a set of intermolecular contacts along the *b* axis and also

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Figure 1

The molecular structure of (I), with the atom-numbering scheme. Displacement ellipsoids are shown at the 50% probability level. H atoms are shown as small circles of arbitrary radii.

contribute to the bonding in the *a*-axis direction. The short intermolecular Br...Br contacts stabilize the structure in the ab plane, as shown in Fig. 3. Along the c-axis direction, the structure is essentially stabilized by interaction between the unsubstituted aromatic ring system and symmetry-related equivalents, also interactions between both Br1 and Br5 with the unsubstituted aromatic ring.

A related packing pattern was found for 2,4-dibromophenyl 4-bromophenyl ether (Eriksson & Hu, 2002), where the bonding in two directions of the unit cell can be described as partly due to $Br \cdot \cdot Br$ interactions. The packing in the third direction is governed mainly by interactions between the aromatic ring systems. The molecules form sheets, with the less substituted ring pointing outwards in every second molecule (cf. Fig. 3) of the title compound, while for the aforementioned 2,4-dibromophenyl 4-bromophenyl ether (Eriksson & Hu, 2002), there are larger cavities where two equivalents of the less substituted ring may fit.

Experimental

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

Crystal data

C ₁₂ H ₅ Br ₅ O	$D_x = 2.582 \text{ Mg m}^{-3}$
$M_r = 564.71$	Mo $K\alpha$ radiation
Monoclinic, $P2_1/c$	Cell parameters from 1573
a = 11.579 (4) Å	reflections
b = 5.3284 (11) Å	$\theta = 1.7-26.0^{\circ}$
c = 24.133 (6) Å	$\mu = 13.82 \text{ mm}^{-1}$
$\beta = 102.66 \ (3)^{\circ}$	T = 293 (2) K
$V = 1452.7 (6) \text{ Å}^3$	Slab, colourless
Z = 4	$0.30 \times 0.09 \times 0.08 \text{ mm}$
Data collection	
Stoe IPDS area-detector	2843 independent reflections
diffractometer	1503 reflections with $I > 2\sigma(I)$
φ scans	$R_{\rm int} = 0.114$
Absorption correction: numerical	$\theta_{\rm max} = 26.0^{\circ}$
(X-RED; Stoe & Cie, 1997)	$h = -14 \rightarrow 14$
$T_{\min} = 0.081, T_{\max} = 0.328$	$k = -6 \rightarrow 6$
18710 measured reflections	$l = -29 \rightarrow 29$



Figure 2

Stereoview of the packing, viewed along the b direction, with the two shortest Br...Br contacts indicated by thick dashed bonds.



Figure 3

Stereoview of the packing, viewed along the c direction, with the two shortest Br...Br contacts indicated by thick dashed bonds. The base plane $(2a \times 3b)$ is marked by the six rectangles.

Refinement

Refinement on F^2 $R[F^2 > 2\sigma(F^2)] = 0.036$ $vR(F^2) = 0.066$	H-atom parameters constrained $w = 1/[\sigma^2(F_o^2) + (0.005P)^2]$ where $P = (F_o^2 + 2F_o^2)/3$
S = 1.03	$(\Delta/\sigma)_{\rm max} < 0.001$
2843 reflections	$\Delta \rho_{\rm max} = 0.55 \ {\rm e} \ {\rm \AA}^{-3}$
63 parameters	$\Delta \rho_{\rm min} = -0.68 \text{ e } \text{\AA}^{-3}$

Table 1

Selected	geometric	parameters	(Å, °)
	D		(

Br1-C2	1.866 (6)	Br5-C6	1.876 (6)
Br2-C3	1.900 (6)	O-C1	1.371 (7)
Br3-C4	1.880 (6)	O-C7	1.403 (8)
Br4-C5	1.882 (6)		
C1-O-C7	118.9 (4)	C12-C7-C8	120.8 (6)
O-C1-C2	119.1 (5)	C12-C7-O	123.8 (6)
O-C1-C6	119.0 (6)	C8-C7-O	115.5 (6)
C2 - C1 - C6	121.6 (6)		
C7-O-C1-C2	100.4 (7)	C1-O-C7-C12	-12.9(9)
C7-O-C1-C6	-86.2 (7)	C1-O-C7-C8	166.8 (5)

Two data sets were recorded with the Stoe IPDS system, using different settings of the crystal. The data sets were brought to a common scale by use of batch scale factors (BASF) determined with *SHELXL*97, and were merged. The rather high internal *R* value of 0.114 decreases to 0.073 when only reflections with $I \ge 2\sigma(I)$ are used.

Data collection: *EXPOSE* in *IPDS* (Stoe & Cie, 1997); cell refinement: *CELL* in *IPDS*; data reduction: *INTEGRATE* in *IPDS* and *X-RED* (Stoe & Cie, 1997); program(s) used to solve structure: *SHELXS*97 (Sheldrick, 1990); program(s) used to refine structure: *SHELXL*97 (Sheldrick, 1997); molecular graphics: *DIAMOND* (Bergerhoff, 1996).

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