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Key indicators

Single-crystal X-ray study T = 174 K Mean σ (C–C) = 0.007 Å Disorder in main residue R factor = 0.038 wR factor = 0.050 Data-to-parameter ratio = 10.5

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

2,2'-Dibromo-3,3',4,4',5,5',6,6'-octamethyl-1,1'-biphenyl

The title compound, $C_{29}H_{24}Br_2$, has crystallographic twofold rotation symmetry. The Br atom in the 2 position is disordered with the methyl group in the 6 position. The biphenyl bridge bond distance and the torsion angle between the rings are in good agreement with similar previously reported structures.

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Comment

The title compound, (I), crystallizes in the space group *Pbcn*, with the molecule on a crystallographic twofold axis, which bisects the angle between the two rings. One half of the molecule consists of a benzene ring, to which are bonded a Br atom and four methyl groups. The Br atom is bonded in the 2 position and is disordered between the 2 and 6 positions. The methyl group in the 6 position is concomitantly disordered.



The major-component C2–Br1 distance is 1.871 (4) Å and the minor-component C6–Br2 distance is 1.809 (5) Å. The major-component C2–Br1 distance is comparable to the C– Br distances in 2,2'-dibromobiphenyl (1.895 and 1.905 Å; MacNeil & Decken, 1999).

The C–C distance of the biphenyl bridge is 1.502 (9) Å, similar to that of 2,2'-dibromobiphenyl (1.499 Å), decachlorobiphenyl (1.522 Å; Pedersen, 1975) and decakis(dichloromethyl)biphenyl (1.534 Å; Biali *et al.*, 1988).

The interplanar angle between the phenyl rings in the molecule is $85.2 (8)^{\circ}$. This compares well with the corresponding angles in 2,2'-dibromobiphenyl (86.0°), decachlorobiphenyl (86.7°) and decakis(dichloromethyl)biphenyl (84.4°).

Experimental

The title compound was synthesized according to a previously reported method (Shibata *et al.*, 2003).

Crystal data

C II D	M. K disting
$C_{20}H_{24}Br_2$	Mo $\kappa \alpha$ radiation
$M_r = 424.22$	Cell parameters from 2260
Orthorhombic, Pbcn	reflections
a = 13.530 (4) Å	$\theta = 2.4-24.0^{\circ}$
b = 10.987 (4) Å	$\mu = 4.59 \text{ mm}^{-1}$
c = 11.916 (3) Å	T = 174.2 K
$V = 1771.4 (9) \text{ Å}^3$	Block, colorless
Z = 4	$0.29 \times 0.21 \times 0.13 \text{ mm}$
$D_x = 1.591 \text{ Mg m}^{-3}$	

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organic papers

Data collection

Bruker SMART 1000	1483 independent reflections
diffractometer	1061 reflections with $I > 3\sigma(I)$
ω scans	$R_{\rm int} = 0.046$
Absorption correction: multi-scan	$\theta_{\rm max} = 24.7^{\circ}$
(Blessing, 1995)	$h = -13 \rightarrow 15$
$T_{\min} = 0.280, T_{\max} = 0.554$	$k = -12 \rightarrow 12$
7776 measured reflections	$l = -13 \rightarrow 13$
Refinement	
Refinement on F	H-atom parameters constraine

H-atom parameters constrained $w = 1/[\sigma^2(F_o) + 0.00022|F_o|^2]$ $(\Delta/\sigma)_{max} < 0.001$ $\Delta\rho_{max} = 0.28 \text{ e } \text{\AA}^{-3}$ $\Delta\rho_{min} = -0.35 \text{ e } \text{\AA}^{-3}$

Table 1

 $\begin{array}{l} R=0.038\\ wR=0.050 \end{array}$

S = 1.85

1061 reflections 101 parameters

Selected geometric parameters (Å, °).

Br1-C2 Br2-C6	1.871 (4) 1.809 (5)	C1-C1 ⁱ	1.502 (9)
$C2 - C1 - C1^{i} - C2^{i}$	-83.7 (8)	$C2 - C1 - C1^i - C6^i$	95.6 (3)

Symmetry code: (i) $1 - x, y, \frac{3}{2} - z$.

Table 2

Contact distances (Å).

$\begin{array}{c} Br1 \cdots Br2^{ii} \\ Br2 \cdots C11^{iii} \end{array}$	3.557 (2) 3.502 (2)	$Br2 \cdots Br2^{iv}$	3.532 (4)
~	1 am	1 43.4	

Symmetry codes: (ii) $x, -y, z - \frac{1}{2}$; (iii) $x, -y, \frac{1}{2} + z$; (iv) 1 - x, -y, 2 - z.

Atoms C10 and C11 were inserted along the C–Br bonds at a fixed distance of 1.54 Å from the ring C atoms. The displacement parameters of these atoms were fixed at 0.0456 Å² because their positions are not well defined and the C–Br distance is too short to allow for proper resolution of the disordered atoms. The Br1 atom occupancy was allowed to refine and then fixed at 0.73. The occupancies of atoms Br2, C10 and C11 were tied appropriately to that of Br1.

Data collection: *SMART* (Bruker, 2003); cell refinement: *SAINT* (Bruker, 2003); data reduction: *SAINT*; program(s) used to solve structure: *SIR*97 (Altomare *et al.*, 1999); program(s) used to refine



Figure 1

The molecular structure of the title compound, including the disorder. Displacement ellipsoids are drawn at the 50% probability level.

structure: *TEXSAN* (Molecular Structure Corporation & Rigaku, 1998); molecular graphics: *TEXSAN*; software used to prepare material for publication: *TEXSAN*.

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