

1,2-Dibromo-4,5-bis(bromomethyl)benzene

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Key indicators

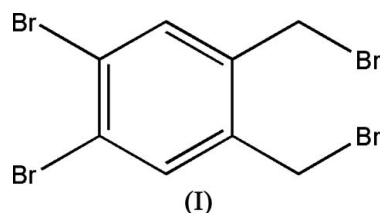
Single-crystal X-ray study
 $T = 292$ K
Mean $\sigma(\text{C}-\text{C}) = 0.007$ Å
 R factor = 0.042
 wR factor = 0.106
Data-to-parameter ratio = 23.7For details of how these key indicators were
automatically derived from the article, see
<http://journals.iucr.org/e>.

In the crystal structure of the title compound, $\text{C}_8\text{H}_6\text{Br}_4$, the packing of the molecules is mainly governed by an intermolecular π - π interaction.

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Comment

Apart from their intrinsic use as dyes and pigments, phthalocyanines show a number of special properties which account for the great research interest related to exciting future prospects (Leznoff & Lever, 1989). Furthermore, phthalocyanine compounds are of considerable interest because of their molecular stacking properties and because of their ability, when doped, to function as electronic conductors (Allcock & Neenan, 1986). The title compound, (I), is an important intermediate for preparation of phthalocyanines (Allcock & Neenan, 1986; Gürek *et al.*, 1991; Gümüş *et al.*, 1992). In this paper, we report the crystal structure of (I) (Fig. 1 and Table 1).



An intermolecular π - π interaction seems to be effective in stabilizing the crystal structure (Fig. 2). The centroid-centroid distance between adjacent benzene rings is 3.786 (3) Å and the interplanar distance is 3.546 (2) Å. There are no inter- or intramolecular hydrogen-bonding interactions.

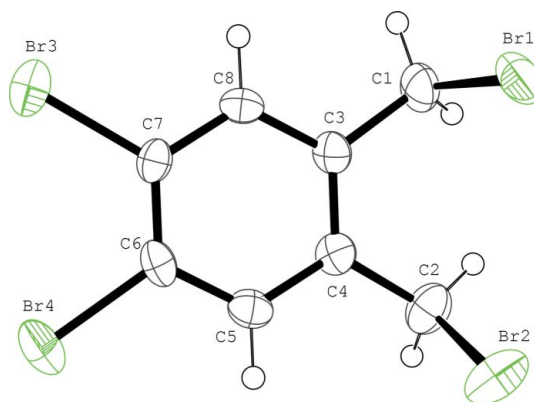


Figure 1
The molecular structure of (I), showing the atom-labeling scheme. Displacement ellipsoids are drawn at the 50% probability level.

Experimental

The title compound was synthesized according to the procedure reported by Allcock & Neenan (1986). Crystals appropriate for X-ray diffraction were obtained by slow evaporation of a petroleum ether solution at 283 K.

Crystal data

$C_8H_6Br_4$ $Z = 8$
 $M_r = 421.77$ $D_x = 2.594 \text{ Mg m}^{-3}$
 Orthorhombic, $Pbca$ Mo $K\alpha$ radiation
 $a = 7.8798 (10) \text{ \AA}$ $\mu = 14.85 \text{ mm}^{-1}$
 $b = 16.225 (2) \text{ \AA}$ $T = 292 (2) \text{ K}$
 $c = 16.894 (2) \text{ \AA}$ Block, colorless
 $V = 2159.8 (5) \text{ \AA}^3$ $0.20 \times 0.20 \times 0.20 \text{ mm}$

Data collection

Bruker SMART 4K CCD area- 2582 independent reflections
 detector diffractometer 1619 reflections with $I > 2\sigma(I)$
 φ and ω scans $R_{int} = 0.081$
 Absorption correction: none $\theta_{max} = 28.0^\circ$
 17417 measured reflections

Refinement

Refinement on F^2 H-atom parameters constrained
 $R[F^2 > 2\sigma(F^2)] = 0.042$ $w = 1/[\sigma^2(F_o^2) + (0.0527P)^2]$
 $wR(F^2) = 0.107$ where $P = (F_o^2 + 2F_c^2)/3$
 $S = 1.00$ $(\Delta/\sigma)_{max} < 0.001$
 2582 reflections $\Delta\rho_{max} = 0.94 \text{ e \AA}^{-3}$
 109 parameters $\Delta\rho_{min} = -1.16 \text{ e \AA}^{-3}$

Table 1

Selected bond lengths (Å).

Br1—C1	1.956 (6)	Br2—C2	1.954 (6)
Br3—C7	1.886 (5)	Br4—C6	1.887 (5)

H atoms were positioned geometrically [$C-H = 0.93$ (CH) and 0.97 \AA (CH_2)] and constrained to ride on their parent atoms [$U_{iso}(H) = 1.2U_{eq}(C)$]. The deepest hole is located 0.72 \AA from atom Br1.

Data collection: SMART (Bruker, 2001); cell refinement: SAINT (Bruker, 2001); data reduction: SAINT; program(s) used to solve structure: SHELXS97 (Sheldrick, 1997); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: PLATON (Spek, 2003); software used to prepare material for publication: SHELXTL (Sheldrick, 2001).

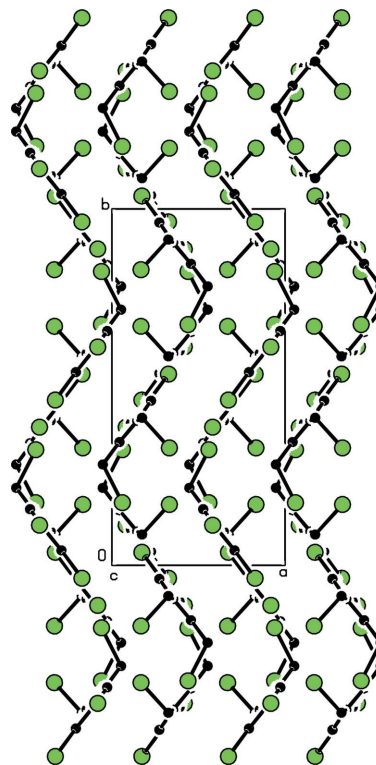


Figure 2

A packing diagram of (I), viewed along the c axis. H atoms have been omitted for clarity.

References

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