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Key indicators

Single-crystal X-ray study T = 297 K Mean σ (C–C) = 0.007 Å R factor = 0.047 wR factor = 0.149 Data-to-parameter ratio = 18.4

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

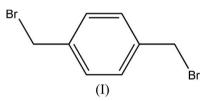
1,4-Bis(bromomethyl)benzene

In the title compound, $C_8H_8Br_2$, the molecules are located on inversion centres and are connected through weak intermolecular $Br \cdots Br$ interactions into layers parallel to the (102) plane.

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Comment

1,4-Bis(halomethyl)benzene compounds have played a very important role in the preparation of bridging ligands used in supramolecular chemistry (Hoskins *et al.*, 1997*a,b*; Wen *et al.*, 2005; Zhang *et al.*, 2005). However, to date only 1,4-bis-(chloromethyl)benzene has been crystallographically characterized (Basaran *et al.*, 1992). In this paper, we report the crystal structure of an isomorphous compound, 1,4-bis-(bromomethyl)benzene, (I).



The asymmetric unit of (I) contains one half-molecule; the molecule is located on an inversion centre (Fig. 1). No unexpected bond lengths or angles are observed.

In the packing of (I), neither hydrogen bonds nor significant $\pi - \pi$ or C-H··· π interactions exist. An analysis of the crystal structure (Spek, 2003) shows that the only possible direction-specific interactions arise from intermolecular Br···Br contacts. In detail, atom Br1 at (x, y, z) interacts with symmetry-related Br1 atoms at $(-x, \frac{1}{2} + y, \frac{1}{2} - z)$ and $(-x, -\frac{1}{2} + y, \frac{1}{2} - z)$, with Br···Br separations of 3.73 (1) Å, slightly

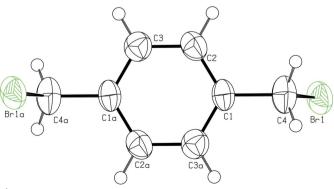


Figure 1

The molecular structure of (I), showing the atom-numbering scheme. Displacement ellipsoids are drawn at the 50% probability level. Atoms labelled with the suffix a are generated by the symmetry operator (1 - x, -y, -z).

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longer than the sum of the van der Waals radii of Br (3.70 Å; Bondi, 1964). As a result, the crystal structure is two-dimensional, parallel to the (102) crystal plane (Fig. 2). No directionspecific interactions are observed between adjacent layers.

Experimental

The title compound was synthesized according to a reported procedure (Hoskins *et al.*, 1997*a*). Crystals suitable for X-ray analysis were formed within 2 d from a methanol solution (0.50 g in 20 ml) at 293 K.

Z = 2

 $D_x = 2.027 \text{ Mg m}^{-3}$

Mo $K\alpha$ radiation

Block, colourless

 $0.30 \times 0.30 \times 0.20$ mm

3213 measured reflections

848 independent reflections

712 reflections with $I > 2\sigma(I)$

 $\mu = 9.30 \text{ mm}^{-1}$

T = 297 (2) K

 $R_{\rm int} = 0.043$

 $\theta_{\rm max} = 26.0^\circ$

Crystal data

 $C_8H_8Br_2$ $M_r = 263.96$ Monoclinic, $P2_1/c$ a = 8.7424 (7) Å b = 4.6275 (4) Å c = 10.8189 (9) Å $\beta = 98.885$ (1)° V = 432.43 (6) Å³

Data collection

Bruker SMART APEX CCD areadetector diffractometer 0.3° wide ω exposures scans Absorption correction: multi-scan (*SADABS*; Sheldrick, 1996) $T_{\min} = 0.082, T_{\max} = 0.157$

Refinement

Refinement on F^2	H-atom parameters constrained
$R[F^2 > 2\sigma(F^2)] = 0.048$	$w = 1/[\sigma^2(F_o^2) + (0.1099P)^2]$
$wR(F^2) = 0.149$	where $P = (F_0^2 + 2F_c^2)/3$
S = 1.06	$(\Delta/\sigma)_{\rm max} < 0.001$
848 reflections	$\Delta \rho_{\rm max} = 0.77 \ {\rm e} \ {\rm \AA}^{-3}$
46 parameters	$\Delta \rho_{\rm min} = -0.85 \text{ e } \text{\AA}^{-3}$

All H atoms were placed in geometrically idealized positions, with C-H = 0.93 (aromatic CH) or 0.97 Å (methylene CH₂). They were included in the refinement in the riding-model approximation, with $U_{iso}(H) = 1.2U_{ea}(C)$.

Data collection: *SMART* (Bruker, 2001); cell refinement: *SMART*; data reduction: *SAINT-Plus* (Bruker, 2001); program(s) used to solve

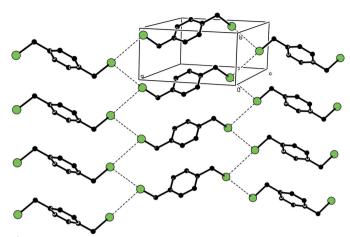


Figure 2

Part of the crystal structure of (I), showing the formation of a layer parallel to (102) through the $Br \cdots Br$ interactions (dashed lines). For the sake of clarity, H atoms have been omitted.

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: *PLATON*.

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