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1,1-Bis(bromomethyl)benzene

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Key indicators

Single-crystal X-ray study $T=298~\mathrm{K}$ Mean $\sigma(\mathrm{C-C})=0.007~\mathrm{Å}$ R factor = 0.050 wR factor = 0.135 Data-to-parameter ratio = 21.2

For 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 molecule, $C_8H_8Br_2$, there is a single $C-H\cdots\pi$ (arene) interaction.

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Comment

The title compound, (I), is used as an organic intermediate in the synthesis of cyclic amines (Chadim *et al.*, 2003) and as a bridging reagent to link two cyclic amine rings (Graham *et al.*, 2005). Its crystal structure is reported here.

$$CH_2Br$$
 CH_2Br
 (I)

The molecular structure of (I) is shown in Fig. 1. The bond lengths and angles are within normal ranges (Allen *et al.*, 1987). Atoms C7 and C8 are almost coplanar with the aryl ring, as reflected in the C7–C1–C2–C3 and C8–C3–C4–C5 torsion angles of –179.1 (4) and –179.5 (5)°, respectively. Atoms Br1 and Br2 are displaced, on opposite sides, by 1.829 (2) and 1.790 (1) Å, respectively, from the least-squares plane defined by atoms C1–C8.

In the crystal structure, there is a weak $C-H\cdots\pi$ (arene) interaction $[H8\cdots Cg^i \ 2.98, \ C8\cdots Cg^i \ 3.550 \ (6) \ Å, \ C8-H8\cdots Cg^i \ 119 \ Å; \ Cg$ is the centroid of the C1-C6 ring; symmetry code: (i) x, -1 + y, z], forming chains propagating along [010] (Fig. 2).

Experimental

The title compound was synthesized by a reported method (Stephenson *et al.*, 1963). Crystals suitable for X-ray analysis were obtained by the slow evaporation of a CH_3CN solution.

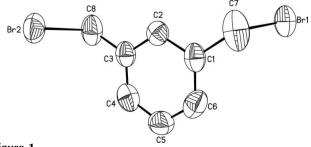


Figure 1
The molecular structure of (I), showing displacement ellipsoids drawn at the 50% probability level. H atoms have been omitted.

© 2006 International Union of Crystallography All rights reserved Crystal data

Data collection

Bruker SMART CCD diffractometer 1933 independent reflections φ and ω scans 1221 reflections with $I > 2\sigma(I)$ Absorption correction: multi-scan (SADABS; Bruker, 2000) $T_{\min} = 0.151, \, T_{\max} = 0.406$ $\theta_{\max} = 27.0^{\circ}$

Refinement

 $\begin{array}{lll} \mbox{Refinement on } F^2 & \mbox{H-atom parameters constrained} \\ R[F^2 > 2\sigma(F^2)] = 0.050 & \mbox{$w = 1/[\sigma^2(F_o^2) + (0.0726P)^2]$} \\ \mbox{$w = (F_o^2) = 0.135$} & \mbox{where } P = (F_o^2 + 2F_o^2)/3$ \\ \mbox{$S = 1.00$} & (\Delta/\sigma)_{\rm max} = 0.001 \\ \mbox{1933 reflections} & \Delta\rho_{\rm max} = 0.75 \ {\rm e \ \mathring{A}^{-3}} \\ \mbox{4 $P_{\rm min} = -0.52$ e \ \mathring{A}^{-3}$} \\ \mbox{$4$ $P_{\rm min} = -0.52$ e \ \mathring$

All H atoms were placed in calculated positions (C—H = 0.93–0.97 Å) and included in the riding-model approximation, with $U_{\rm iso}({\rm H})$ = $1.2 U_{\rm eq}({\rm C})$.

Data collection: *SMART* (Bruker, 2000); cell refinement: *SAINT* (Bruker, 2000); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *SHELXTL* (Bruker, 2000); software used to prepare material for publication: *SHELXTL*.

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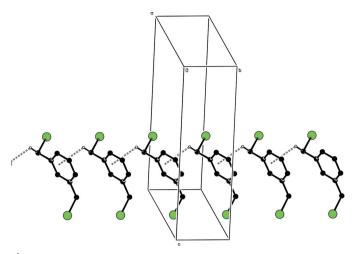


Figure 2 Part of the crystal structure of (I), showing $C-H\cdots\pi$ (arene) interactions as dashed lines. For clarity, H atoms not in the motif shown have been omitted

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