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Key indicators

Single-crystal X-ray study T = 120 KMean σ (C–C) = 0.003 Å R factor = 0.056 wR factor = 0.166 Data-to-parameter ratio = 13.4

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

Hexakis(hydroxymethyl)benzene

The centrosymmetric title compound, $C_{12}H_{18}O_6$, exhibits a three-dimensional network structure. The structure displays extensive $O-H\cdots O$ hydrogen bonding. In each molecule, two sets of three neighboring hydroxy groups are located above and below the plane defined by the central benzene ring.

Comment

The title compound, (I) (Fig. 1), forms a hydrogen-bonded three-dimensional network in the solid state (Table 1). Figs. 2 and 3 illustrate the packing viewed along the a and b axes.



Experimental

The title compound was prepared according to a published procedure and recrystallized from methanol (Alphonse *et al.*, 1988).





Figure 1

© 2007 International Union of Crystallography All rights reserved The molecular structure of (I) (displacement ellipsoids are at the 50% probability level). H atoms are drawn as spheres of arbitrary radius.

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Data collection

Stoe IPDSII diffractometer Absorption correction: none 4212 measured reflections

Refinement

 $R[F^2 > 2\sigma(F^2)] = 0.056$ 85 parameters $wR(F^2) = 0.166$ H-atom parameters constrainedS = 0.96 $\Delta \rho_{max} = 0.31$ e Å⁻³1143 reflections $\Delta \rho_{min} = -0.33$ e Å⁻³

Table 1

Hydrogen-bond geometry (Å, °).

D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - H \cdot \cdot \cdot A$
0.84	2.34	2.951 (2)	130
0.84	1.92	2.739 (2)	165
0.84	2.71	3.049 (3)	106
0.84	2.18	2.906 (3)	145
0.84	1.95	2.767 (2)	163
	<i>D</i> —H 0.84 0.84 0.84 0.84 0.84 0.84	$\begin{array}{c cccc} D-H & H\cdots A \\ \hline 0.84 & 2.34 \\ 0.84 & 1.92 \\ 0.84 & 2.71 \\ 0.84 & 2.18 \\ 0.84 & 1.95 \\ \end{array}$	$\begin{array}{c ccccccccccccccccccccccccccccccccccc$

1143 independent reflections

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

 $R_{\rm int} = 0.088$

Symmetry codes: (i) $-x, y + \frac{1}{2}, -z + \frac{1}{2}$; (ii) $x, -y + \frac{3}{2}, z + \frac{1}{2}$; (iii) $x - 1, -y + \frac{3}{2}, z - \frac{1}{2}$.

H atoms were treated as riding, with C-H = 0.99 Å, O-H = 0.84 Å and $U_{iso}(H)$ values set at $1.2U_{eq}(C)$ and $1.5U_{eq}(O)$.

Data collection: X-AREA (Stoe & Cie, 2002); cell refinement: X-AREA; data reduction: X-AREA; program(s) used to solve structure: SHELXS97 (Sheldrick, 1997a); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997a); molecular graphics: SHELXTL (Sheldrick, 1997b); software used to prepare material for publication: SHELXTL.

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References

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- Sheldrick, G. M. (1997a). SHELXS97 and SHELXL97. University of Göttingen, Germany.
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Figure 2

Packing diagram of (I) (viewed along the a axis). Hydrogen bonding is indicated by dashed lines.



Figure 3

Packing diagram of 1 (viewed along the b axis). Hydrogen bonding is indicated by dashed lines.

Stoe & Cie (2002). *X-AREA* (Version 1.18) and *X-RED32* (Version 1.04). Stoe & Cie, Darmstadt, Germany.