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Key indicators

Single-crystal X-ray study
 T = 173 K
 Mean σ (C–C) = 0.004 Å
 Disorder in main residue
 R factor = 0.057
 wR factor = 0.213
 Data-to-parameter ratio = 22.4

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

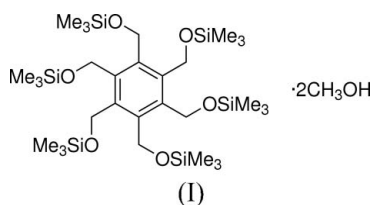
Hexakis(trimethylsiloxymethyl)benzene methanol disolvate

In the structure of the centrosymmetric title compound, $C_{30}H_{66}O_6Si_6 \cdot 2CH_3OH$, OSiMe₃ groups are located above and below the plane defined by the central benzene ring in an alternating fashion.

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Comment

The molecular structure of $C_{32}H_{74}O_8Si_6$, (I), is shown in Fig. 1. The benzene ring is located on an inversion centre. Atom O1 is disordered over two positions (1:1). One methanol solvent molecule is disordered over two positions (3:1).



Experimental

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

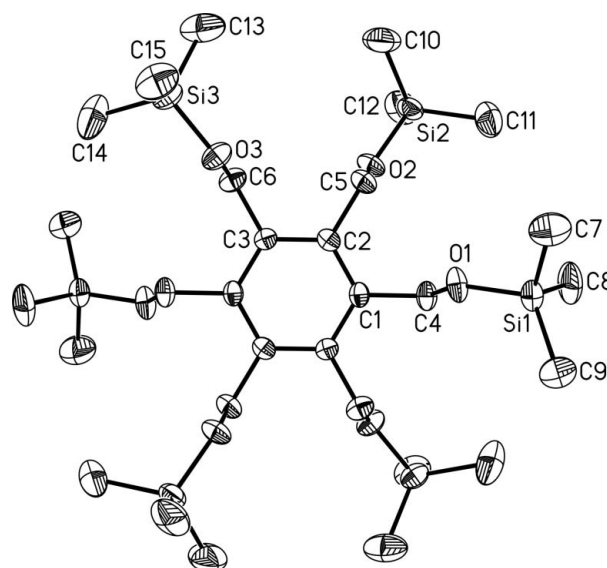


Figure 1
 The molecular structure of (I) (displacement ellipsoids are drawn at the 50% probability level). H atoms are drawn as spheres of arbitrary radius. Only one disordered component is shown; methanol solvent molecules have been omitted for clarity. Unlabelled atoms are related to labelled atoms by $(1 - x, 1 - y, 1 - z)$.

Crystal data

C₃₀H₆₆O₆Si₆·2CH₄OM_r = 755.45Triclinic, P $\bar{1}$

a = 9.7075 (19) Å

b = 10.153 (2) Å

c = 12.227 (2) Å

 α = 101.34 (3)° β = 97.61 (4)° γ = 96.04 (3)°V = 1160.4 (4) Å³

Z = 1

Mo K α radiation μ = 0.22 mm⁻¹

T = 173 (2) K

0.30 × 0.30 × 0.20 mm

Data collection

Stoe Stadi-4 diffractometer

Absorption correction: none

9084 measured reflections

5117 independent reflections

3168 reflections with $I > 2\sigma(I)$ $R_{\text{int}} = 0.039$

Refinement

 $R[F^2 > 2\sigma(F^2)] = 0.057$ $wR(F^2) = 0.213$

S = 1.07

5117 reflections

228 parameters

1 restraint

H-atom parameters constrained

 $\Delta\rho_{\text{max}} = 0.77 \text{ e \AA}^{-3}$ $\Delta\rho_{\text{min}} = -1.06 \text{ e \AA}^{-3}$

Table 1

Hydrogen-bond geometry (Å, °).

D—H...A	D—H	H...A	D...A	D—H...A
O4'—H4'...O1	0.84	2.41	2.981 (15)	126
O4'—H4'...O1'	0.84	2.73	3.382 (15)	135
O4—H4...O3	0.84	2.59	2.863 (6)	101

H atoms were treated as riding, with C—H = 0.98–0.99 Å, O—H = 0.84 Å, and $U_{\text{iso}}(\text{H})$ values set at 1.2 or 1.5 times $U_{\text{eq}}(\text{C})$ and 1.5 times $U_{\text{eq}}(\text{O})$. The methanol solvent molecule was refined using isotropic displacement parameters and one restraint to fix a C—O bond distance. The methanol molecule is disordered over two positions. One CH₂—O group of the main molecule is disordered equally over two positions.

Data collection: *STADIA* (Stoe & Cie, 1995); cell refinement: *STADIA*; data reduction: *STADIA*; 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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