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Key indicators

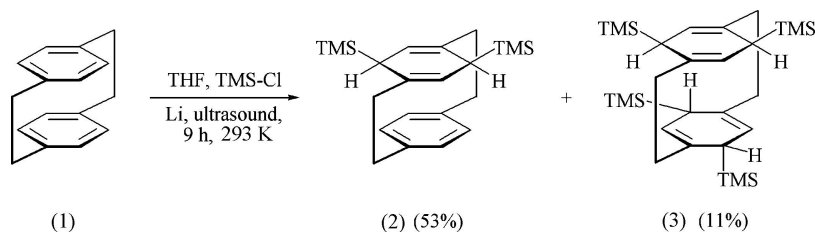
Single-crystal X-ray study
 $T = 143$ K
Mean σ (Wae) = 0.004 Å
 R factor = 0.040
 wR factor = 0.114
Data-to-parameter ratio = 14.6For details of how these key indicators were
automatically derived from the article, see
<http://journals.iucr.org/e>.***syn-exo-4,7,12,15-Tetrakis(trimethylsilyl)-***
4,7,12,15-tetrahydro[2.2]paracyclophaneThe title compound, $C_{28}H_{52}Si_4$, crystallizes in the uncommon
space group $I4_122$ with 222 symmetry. Despite the reduction of
the rings, several features of [2.2]paracyclophane strain are
still apparent.

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Comment

In the previous paper (Jones *et al.*, 2005), in which intro-
ductory material can be found, we have presented the struc-
ture of compound (2) in the scheme. We present here the
structure of the doubly reduced compound (3), which was
formed in the same reaction.The structure of (3) is shown in Fig. 1. It crystallizes in the
space group $I4_122$, which is uncommon; a search of the
Cambridge Structural Database (Version 5.26; Allen, 2002)
revealed only 23 other examples. The molecule displays
crystallographic symmetry 222, and the numbering of the
asymmetric unit therefore cannot conform to the IUPAC
numbering of the whole molecule.The reduced rings display flattened and somewhat distorted
twist conformations with significantly non-zero torsion angles
about the double bonds $C2=C4$. The molecule, despite the
reduction of both rings, still shows features of strain that
are typical of [2.2]paracyclophane systems, *e.g.* the bridgehead
distance $C2 \cdots C2^i = 2.816$ (5) Å [symmetry code: (i) $\frac{1}{2} - y, \frac{1}{2} - x, \frac{3}{2} - z$] the lengthened single bonds and widened angles in
the bridges.

Experimental

The title compound was prepared as described by Hopf *et al.* (1997).
Single crystals were grown by evaporation of a solution in *n*-hexane.

Crystal data

 $C_{28}H_{52}Si_4$
 $M_r = 501.06$
Tetragonal, $I4_122$
 $a = 13.906$ (4) Å
 $c = 16.263$ (5) Å
 $V = 3144.9$ (16) Å³
 $Z = 4$
 $D_x = 1.058$ Mg m⁻³Mo $K\alpha$ radiation
Cell parameters from 66
reflections
 $\theta = 10\text{--}11.5^\circ$
 $\mu = 0.20$ mm⁻¹
 $T = 143$ (2) K
Block, colourless
 $0.6 \times 0.6 \times 0.5$ mm

Data collection

Stoe Stadi-4 diffractometer
 ω/θ scans
 Absorption correction: none
 3911 measured reflections
 1068 independent reflections
 936 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.031$

$\theta_{\text{max}} = 27.5^\circ$
 $h = 0 \rightarrow 18$
 $k = -18 \rightarrow 0$
 $l = -21 \rightarrow 21$
 3 standard reflections
 frequency: 60 min
 intensity decay: none

Refinement

Refinement on F^2
 $R[F^2 > 2\sigma(F^2)] = 0.040$
 $wR(F^2) = 0.114$
 $S = 1.07$
 1068 reflections
 73 parameters
 H-atom parameters constrained

$w = 1/[\sigma^2(F_o^2) + (0.0668P)^2 + 1.0612P]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\text{max}} < 0.001$
 $\Delta\rho_{\text{max}} = 0.23 \text{ e } \text{Å}^{-3}$
 $\Delta\rho_{\text{min}} = -0.14 \text{ e } \text{Å}^{-3}$

Table 1

Selected geometric parameters (Å, °).

C1—C1 ⁱ	1.571 (5)	C2—C3	1.506 (4)
C2—C4	1.326 (4)		
C2—C1—C1 ⁱ	112.36 (15)	C2—C3—C4	109.7 (2)
C4—C2—C3	120.2 (2)	C2—C4—C3	124.8 (2)
C4—C2—C3—C4	28.7 (3)	C2—C3—C4—C2	−18.9 (3)
C3—C2—C4—C3	−10.2 (4)		

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

Methyl H atoms were placed in ideally staggered positions and then refined using a riding model, with C—H 0.98 Å and H—C—H = 109.5°. Other H atoms were included using a riding model, with C—H = 0.95 (sp^2), 0.99 (CH_2) or 1.00 Å (methine). Refinement with Friedel opposite reflections treated as independent led to a Flack (1983) parameter insignificantly different from 0.5 (indicating inversion twinning) and Friedel opposite reflections were therefore merged for the final refinement. To improve stability of refinement in view of the moderate data/parameter ratio, anisotropic displacement parameters were subject to similarity and rigid-bond restraints.

Data collection: *DIF4* (Stoe & Cie, 1992); cell refinement: *DIF4*; data reduction: *REDU4* (Stoe & Cie, 1992); program(s) used to solve structure: *SHELXS97* (Sheldrick, 1990); program(s) used to refine

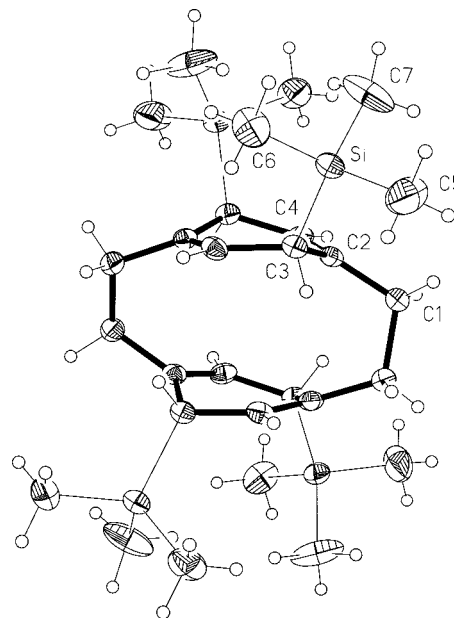


Figure 1

The molecule of compound (3) in the crystal structure. Displacement ellipsoids are drawn at the 30% probability level.

structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *XP* (Siemens, 1994); software used to prepare material for publication: *SHELXL97*.

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