Received 20 April 2006

Accepted 24 April 2006

Acta Crystallographica Section E Structure Reports Online

ISSN 1600-5368

Ralf Hänel,^a Henning Hopf^a and Peter G. Jones^b*

^aInstitut für Organische Chemie, Technische Universität Braunschweig, Postfach 3329, 38023 Braunschweig, Germany, and ^bInstitut für Anorganische und Analytische Chemie, Technische Universität Braunschweig, Postfach 3329, 38023 Braunschweig, Germany

Correspondence e-mail: p.jones@tu-bs.de

Key indicators

Single-crystal X-ray study T = 173 K Mean σ (C–C) = 0.003 Å R factor = 0.056 wR factor = 0.148 Data-to-parameter ratio = 16.8

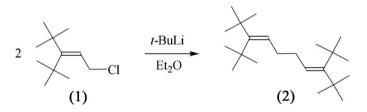
For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e.

3,8-Di-*tert*-butyl-2,2,9,9-tetramethyldeca-3,7-diene

The title compound, $C_{22}H_{42}$, displays crystallographic inversion symmetry. The conformation of the central chain is defined by the torsion angle C=C-C-C of 124.3 (3)°. The C=C-C angle in the chain is 133.2 (2)°. The central C-C bond is short at 1.516 (4) Å.

Comment

During our studies on highly *tert*-butylated hydrocarbons (Hänel, 1996; Hopf *et al.*, 2001) we investigated the behaviour of the allyl chloride (1) towards various metals and metal-organic reagents. When (1) is reacted with *tert*-butyl lithium in diethyl ether at 203 K it dimerizes and yields the diene (2). Here we present the structure of (2).



The molecule of (2), which displays crystallographic inversion symmetry, is shown in Fig. 1. Some evidence of steric crowding is provided by short intramolecular H...H contacts, such as $H1A\cdots H11C = 1.82$ Å, $H1A\cdots H9A = 1.81$ Å and $H5A \cdots H7A = 1.85$ Å; there are no unusually short intermolecular contacts. The conformation of the central chain is largely defined by the $C3=C4-C5-C5^{i}$ torsion angle (Table 1); the torsion angle around the central bond $C5-C5^{i}$ is 180° by symmetry. Molecular dimensions (Table 1) may be regarded as normal, such as the C3=C4 and C4-C5 bond lengths and the C3=C4-C5 angle. A similar C=C bond length, 1.349 (2) Å, and C = C - C angle, 131.1 (2)°, were observed in 1,1,4,4-tetra-tert-butyl-1,3-butadiene [(3); Hopf et al., 1994, 1996). The $C5-C5^{i}$ bond seems rather short at 1.516 (4) A; no other *tert*-butyl-substituted chain of a similar nature is available for comparison. Some caution is indicated in view of problems with the structure of (3), in which contamination by small quantities of an impurity initially led to systematically incorrect dimensions, later corrected (Hopf et al., 1994, 1996).

Experimental

To a solution of (1) (1.0 g, 5.3 mmol) in diethyl ether (7 ml) was added over a period of 30 min at 203 K a 1.7 M tert-butyllithium solution (6.3 ml) in diethyl ether. The mixture was stirred for 30 min at room temperature and refluxed for 12 h to complete the

All rights reserved

© 2006 International Union of Crystallography

dimerization. After work-up, solvent removal and recrystallization (ethanol), (2) (280 mg, 34%) was obtained. The spectroscopic data agree with the proposed structure (Hänel, 1996).

 $V = 510.0 (5) \text{ Å}^3$ Z = 1

 $D_x = 0.998 \text{ Mg m}^{-3}$

Mo $K\alpha$ radiation $\mu = 0.06 \text{ mm}^{-1}$

Tablet, colourless

 $0.7 \times 0.5 \times 0.2 \ \text{mm}$

3 standard reflections

frequency: 60 min

intensity decay: none

 $w = 1/[\sigma^2(F_0^2) + (0.0574P)^2]$

where $P = (F_0^2 + 2F_c^2)/3$

+ 0.1511P]

 $(\Delta/\sigma)_{\rm max} < 0.001$

 $\Delta \rho_{\rm max} = 0.16 \text{ e} \text{ Å}^{-3}$

 $\Delta \rho_{\rm min} = -0.17 \text{ e} \text{ Å}^{-3}$

T = 173 (2) K

 $R_{\rm int} = 0.057$

 $\theta_{\rm max} = 24.9^\circ$

Crystal data

 $\begin{array}{l} C_{22}H_{42} \\ M_r = 306.56 \\ \text{Triclinic, } P\overline{1} \\ a = 6.350 \ (4) \ \mathring{A} \\ b = 8.161 \ (4) \ \mathring{A} \\ c = 10.655 \ (6) \ \mathring{A} \\ \alpha = 87.47 \ (4)^{\circ} \\ \beta = 88.08 \ (4)^{\circ} \\ \gamma = 67.63 \ (4)^{\circ} \end{array}$

Data collection

Stoe STADI-4 diffractometer ω/θ scans Absorption correction: none 2714 measured reflections 1783 independent reflections 1201 reflections with $I > 2\sigma(I)$

Refinement

Refinement on F^2 $R[F^2 > 2\sigma(F^2)] = 0.056$ $wR(F^2) = 0.148$ S = 1.081783 reflections 106 parameters H-atom parameters constrained

Table 1

Selected geometric parameters (Å, °).

C3-C4 C4-C5	1.336 (3) 1.508 (3)	C5-C5 ⁱ	1.516 (4)
C3-C4-C5	133.2 (2)	C4-C5-C5 ⁱ	112.0 (2)
C3-C4-C5-C5 ⁱ	124.3 (3)		
Symmetry code: (i) $-x +$	1, -y, -z.		

Methyl H atoms were located in difference syntheses, idealized $(C-H = 0.98 \text{ Å} \text{ and } H-C-H = 110^{\circ})$ and refined on the basis of

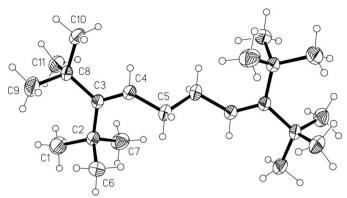


Figure 1

The molecular structure of the title compound. Displacement ellipsoids are drawn at the 50% probability level.

rigid groups allowed to rotate but not tip. Other H atoms were placed in calculated positions (C–H = 0.95 Å for sp^2 CH and 0.99 Å for CH₂) and allowed to ride on their parent C atoms. All U_{iso} (H) values were fixed at 1.2 times U_{eq} of the parent atom.

Data collection: *DIF4* (Stoe & Cie, 1992); cell refinement: *DIF4*; data reduction: *REDU4* (Stoe & Cie, 1992); program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *XP* (Siemens, 1994); software used to prepare material for publication: *SHELXL97*.

We thank Mr A. Weinkauf for technical assistance.

References

Hänel, R. (1996). PhD thesis, Technical University of Braunschweig, Germany.

Hopf, H., Hänel, R., Jones, P. G. & Bubenitschek, P. (1994). Angew. Chem. Int. Ed. Engl. 33, 1369–1370.

- Hopf, H., Hänel, R., Jones, P. G. & Bubenitschek, P. (1996). Angew. Chem. Int. Ed. Engl. 35, 222.
- Hopf, H., Mlynek, C., Klein, D., Traetteberg, M. & Bakken, P. (2001). Eur. J. Org. Chem. pp. 1385–1391.

Sheldrick, G. M. (1997). SHELXL97 and SHELXS97. University of Göttingen, Germany.

Siemens (1994). XP. Version 5.03. Siemens Analytical X-ray Instruments Inc., Madison, Wisconsin, USA.

Stoe & Cie (1992). DIF4 and REDU4. Stoe & Cie, Darmstadt, Germany.