Acta Crystallographica Section E

Structure Reports Online

ISSN 1600-5368

Christian Näther, a* Oliver Krüger and Uta Wille

^aInstitut für Anorganische Chemie, Christian-Albrechts-Universität Kiel, Olshausenstraße 40, D-24098 Kiel, Germany, and ^bInstitut für Organische Chemie, Christian-Albrechts-Universität Kiel, Olshausenstraße 40, D-24098 Kiel, Germany

Correspondence e-mail: cnaether@ac.uni-kiel.de

Key indicators

Single-crystal X-ray study T = 170 KMean $\sigma(\text{C-C}) = 0.002 \text{ Å}$ R factor = 0.049 wR factor = 0.140Data-to-parameter ratio = 17.8

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

cis-[4a]-cisoid-[4a,4b]-cis-[4b]-8a,8b-Diethyl-1,3,6,8-tetramethylperhydro-1,3,6,8-tetraazabiphenylene-2,4,5,7-tetraone

The title compound, $C_{16}H_{24}N_4O_4$, was prepared by [2+2]-photocycloaddition of 1,3-dimethyl-6-ethyluracil in acetone. The structure determination was undertaken in order to determine the stereoconfiguration of the product which could not be extracted from NMR data. Single-crystal X-ray analysis revealed that the methyl groups are located on the same side of the cyclobutane ring (cis) and that the monomers are oriented head-to-head (syn). There are two crystallographically independent molecules in the asymmetric unit; these differ only slightly in geometry.

Received 27 February 2002 Accepted 8 March 2002 Online 15 March 2002

Experimental

The title compound was prepared by [2+2]-photocycloaddition of 1,3-dimethyl-6-ethyluracil in acetone. The products were separated by column chromatography. Single crystals were obtained by slow evaporation of n-pentane into a saturated solution of the title compound in dichloromethane. Details of the synthesis are given by Krüger (2002).

Crystal data

 $C_{16}H_{24}N_4O_4$ Z = 4 $D_x = 1.349 \text{ Mg m}^{-3}$ $M_r = 336.39$ Triclinic, $P\overline{1}$ Mo $K\alpha$ radiation a = 8.5284 (6) Å Cell parameters from 8000 b = 12.2178 (9) Åreflections c = 16.2102 (12) Å $\theta = 10.5 - 27^{\circ}$ $\mu = 0.10 \text{ mm}^{-1}$ $\alpha = 91.481 (9)^{\circ}$ $\beta = 101.269 (8)^{\circ}$ T = 170 (2) K $\gamma = 90.518 (8)^{\circ}$ Needle, colourless $V = 1655.8 (2) \text{ Å}^3$ $0.40 \times 0.10 \times 0.08 \text{ mm}$

Data collection

 $\begin{array}{lll} \text{Stoe Imaging Plate Diffraction} & 5552 \text{ reflections with } I > 2\sigma(I) \\ \text{System diffractometer} & R_{\text{int}} = 0.033 \\ \varphi \text{ scans} & \theta_{\text{max}} = 28.1^{\circ} \\ \text{Absorption correction: none} & h = -11 \rightarrow 11 \\ 15745 \text{ measured reflections} & k = -16 \rightarrow 16 \\ 7846 \text{ independent reflections} & l = -21 \rightarrow 20 \\ \end{array}$

Refinement

Refinement on F^2 $w = 1/[\sigma^2(F_o^2) + (0.0822P)^2]$ $R[F^2 > 2\sigma(F^2)] = 0.049$ + 0.1459P] where $P = (F_o^2 + 2F_c^2)/3$ S = 1.02 $(\Delta/\sigma)_{max} = 0.001$ $\Delta\rho_{max} = 0.26$ e Å $^{-3}$ 442 parameters $\Delta\rho_{min} = -0.31$ e Å $^{-3}$ Extinction correction: SHELXL97 Extinction coefficient: 0.035 (2)

© 2002 International Union of Crystallography Printed in Great Britain – all rights reserved

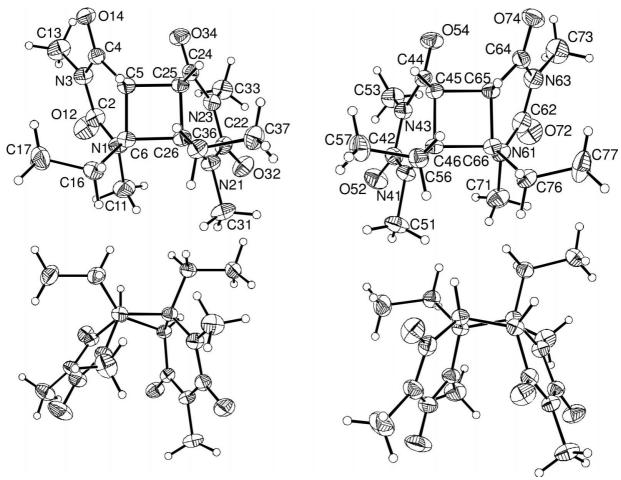


Figure 1
Top view (top) and side view (bottom) of the structures of the two independent molecules of the title compound, with atom labelling and displacement ellipsoids drawn at the 50% probability level.

The methine and methylene H atoms were positioned with idealized geometry (C— $H_{\rm methine}=1.00~{\rm \mathring{A}}$ and C— $H_{\rm methylene}=0.99~{\rm \mathring{A}}$). The positions of the methyl H atoms were idealized (C—H = 0.98 Å), then refined as rigid groups allowed to rotate but not tip. All H atoms were refined with fixed isotropic displacement parameters using a riding model with $U_{\rm iso}=1.2U_{\rm eq}({\rm C})$ and $1.5U_{\rm eq}({\rm C})$ for methine/methylene and methyl H atoms, respectively.

Data collection: *IPDS Program Package* (Stoe & Cie, 1998); cell refinement: *IPDS Program Package*; data reduction: *IPDS Program Package*; program(s) used to solve structure: *SHELXS*97 (Sheldrick, 1997); program(s) used to refine structure: *SHELXL*97 (Sheldrick, 1997); molecular graphics: *XP* in *SHELXTL* (Bruker, 1998); software used to prepare material for publication: *CIFTAB* in *SHELXL*97.

This work is supported by the state of Schleswig-Holstein and the Deutsche Forschungsgemeinschaft. We are very thankful to Professor Dr Wolfgang Bensch for the opportunity to use his experimental equipment.

References

Bruker (1998). SHELXTL. Version 5.1. Bruker AXS Inc., Madison, Wisconsin, USA.

Krüger, O. (2002). PhD Thesis, University of Kiel, Germany.

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

Stoe & Cie (1998). IPDS Program Package. Version 2.89. Stoe & Cie, Darmstadt, Germany.