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Key indicators

Single-crystal X-ray study T = 170 KMean $\sigma(\text{C-C}) = 0.002 \text{ Å}$ R factor = 0.042 wR factor = 0.121Data-to-parameter ratio = 17.0

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

cis-[4a]-transoid-[4a,4b]-cis-[4b]-1,3,6,8,8a,8b-Hexa-methylperhydro-1,3,6,8-tetraazabiphenylene-2,4,5,7-tetraone

The title compound, $C_{14}H_{20}N_4O_4$, was prepared by [2+2]-photocycloaddition of 1,3,6-trimethyluracil 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 opposite sides of the cyclobutane ring (*trans*) and that the monomers are oriented head-to-head (*syn*).

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Experimental

The title compound was prepared by [2+2]-photocycloaddition of 1,3,6-trimethyluracil 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

 $C_{14}H_{20}N_4O_4$ $M_r = 308.34$ $D_x = 1.401 \text{ Mg m}^{-3}$ Triclinic, $P\overline{1}$ Mo $K\alpha$ radiation a = 7.6398 (6) Å Cell parameters from 8000 b = 8.6267 (7) Åreflections c = 12.308 (1) Å $\theta = 10.5 – 27^{\circ}$ $\mu = 0.10 \text{ mm}^{-1}$ $\alpha = 71.483 (9)^{\circ}$ $\beta = 80.13 (1)^{\circ}$ T = 170 (2) KIrregular polyhedron, colourless $\gamma = 72.521 (9)^{\circ}$ $V = 731.09 (10) \text{ Å}^3$ $0.4 \times 0.2 \times 0.2 \text{ mm}$

Data collection

Refinement

 $\begin{array}{lll} \text{Stoe Imaging Plate Diffraction} & 3019 \text{ reflections with } I > 2\sigma(I) \\ \text{System diffractometer} & R_{\text{int}} = 0.035 \\ \varphi \text{ scans} & \theta_{\text{max}} = 28.1^{\circ} \\ \text{Absorption correction: none} & h = -10 \rightarrow 10 \\ 8266 \text{ measured reflections} & k = -11 \rightarrow 11 \\ 3499 \text{ independent reflections} & l = -16 \rightarrow 16 \\ \end{array}$

 $\begin{array}{lll} \text{Refinement on } F^2 & w = 1/[\sigma^2(F_o^2) + (0.0731P)^2 \\ R[F^2 > 2\sigma(F^2)] = 0.042 & + 0.1861P] \\ wR(F^2) = 0.121 & \text{where } P = (F_o^2 + 2F_c^2)/3 \\ S = 1.05 & (\Delta/\sigma)_{\text{max}} < 0.001 \\ 3499 \text{ reflections} & \Delta\rho_{\text{max}} = 0.39 \text{ e} \mathring{A}^{-3} \\ 206 \text{ parameters} & \Delta\rho_{\text{min}} = -0.31 \text{ e} \mathring{A}^{-3} \\ \text{H-atom parameters constrained} & \text{Extinction correction: } SHELXL97 \end{array}$

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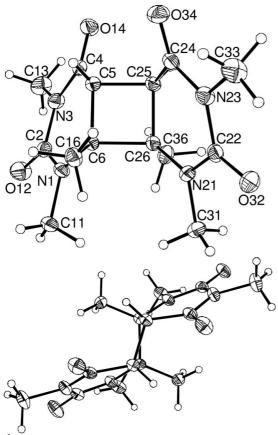


Figure 1
Top view (top) and side view (bottom) of the molecular structure of the title compound, with atom labelling and displacement ellipsoids drawn at the 50% probability level.

Extine methine H. atoms (were positioned with idealized geometry (C— $H_{\rm methine} = 1.00$ Å). 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 $U_{\rm iso} = 1.2 U_{\rm eq}(C)$ and $1.5 U_{\rm eq}(C)$ for methine 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.

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