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Key indicators

Single-crystal X-ray study T = 170 KMean $\sigma(C-C) = 0.002 \text{ Å}$ R factor = 0.040wR factor = 0.111 Data-to-parameter ratio = 18.6

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]-4b-Ethyl-1,3,5,7,8bhexamethylperhydro-1,3,5,7-tetraazabiphenylene-2,4,6,8-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 and the ethyl group are located on the same side of the cyclobutane ring (cis) and that the monomers are oriented head-to-tail (anti).

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Experimental

The title compound was formed as a byproduct in the [2+2]-photocycloaddition of 1,3-dimethyl-6-ethyluracil in acetone, by a mechanism which is not yet clear. 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

 $C_{16}H_{24}N_4O_4$ Mo $K\alpha$ radiation $M_{\rm r} = 336.39$ Cell parameters from 8000 Orthorhombic, Pbca reflections a = 15.2661 (9) Å $\theta = 17-23.5^{\circ}$ $\mu = 0.09 \text{ mm}^{-1}$ b = 14.4367 (8) Åc = 15.7200 (12) ÅT = 170 (2) K $V = 3464.6 (4) \text{ Å}^3$ Irregular block, colourless Z = 8 $0.5 \times 0.3 \times 0.3 \text{ mm}$ $D_x = 1.290 \text{ Mg m}^{-3}$

Data collection

Stoe Imaging Plate Diffraction 3642 reflections with $I > 2\sigma(I)$ $R_{\rm int} = 0.038$ System diffractometer $\theta_{\text{max}} = 28.1^{\circ}$ $h = -20 \rightarrow 20$ Absorption correction: none $k=-17\to18$ 27222 measured reflections $l = -20 \rightarrow 20$ 4171 independent reflections

Refinement

Refinement on F^2 $w = 1/[\sigma^2(F_o^2) + (0.0617P)^2]$ $R[F^2 > 2\sigma(F^2)] = 0.040$ + 0.9107P $wR(F^2) = 0.111$ where $P = (F_o^2 + 2F_c^2)/3$ S = 1.04 $(\Delta/\sigma)_{\text{max}} < 0.001$ $\Delta \rho_{\text{max}} = 0.001$ $\Delta \rho_{\text{max}} = 0.29 \text{ e Å}^{-3}$ $\Delta \rho_{\text{min}} = -0.19 \text{ e Å}^{-3}$ 4171 reflections 224 parameters Extinction correction: SHELXL97 H-atom parameters constrained Extinction coefficient: 0.0088 (11)

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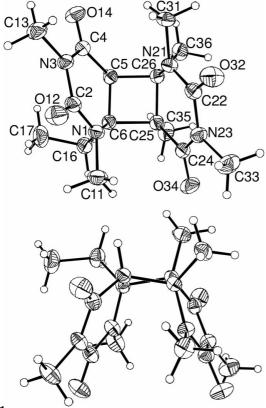


Figure 1
Top view (top) and side view (bottom) of the molecular structure of the title compound, with atom labelling and displacement ellipsoids drawn at the 50% probability level.

The methine and methylene H atoms were positioned with idealized geometry (C-H_{methine} = 1.00 Å and C-H_{methylene} = 0.99 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/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.

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