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Key indicators

Single-crystal X-ray study T = 292 KMean σ (C–C) = 0.003 Å R factor = 0.049 wR factor = 0.148 Data-to-parameter ratio = 14.2

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

Dimethyl 2,6-di-*tert*-butyl-4,8-dioxo-1,2,5,6tetrahydro-2,3a,4a,6,7a,8a-hexaazacyclopenta-[*def*]fluorene-8b,8c-dicarboxylate

The title compound, $C_{20}H_{32}N_6O_6$, is a glycoluril derivative. The molecule is built up from four fused rings, namely two nearly planar imidazole five-membered rings and two non-planar triazine six-membered rings. In the five-membered imidazole rings, the N-C_{carbonyl} bond distances are much shorter than the other N-C distances, indicating π - π conjugation within the nearly planar ring. Both six-membered rings display chair conformations.

Comment

Glycoluril derivatives have applications in many fields, such as explosives, slow-release fertilizers, crosslinkers, iodogens, stabilisers of organic compounds against photodegradation and reagents in combinatorial chemistry (Wu *et al.*, 2002). As part of our ongoing investigation of glycoluril derivatives (Li & Wu, 2005), we present here the structure of the title compound, (I).



The molecular structure of (I) is shown in Fig. 1. The molecule is built up from four fused rings, namely two nearly planar imidazole five-membered rings and two non-planar triazine six-membered rings. Within the nearly planar five-membered ring, the N-C_{carbonyl} bond distances are much shorter than the other N-C distances (Table 1). This is due to the occurrence of π - π conjugation.

Experimental

tert-Butylamine (0.73 g, 10 mmol) and formaldehyde (1.2 g, 40 mmol) were added to a stirred solution of 2,5-dioxotetrahydroimidazo[4,5-d]imidazole-3a,6a-dicarboxylic acid dimethyl ester (1.29 g, 5 mmol) in acetonitrile (50 ml) under a nitrogen atmosphere. The mixture was stirred overnight at room temperature. The solvent was evaporated to dryness and the compound was purified by column-chromatography to yield (I) (2.03 g, 90%) as a colourless solid. Colourless block-like crystals of (I) suitable for data collection were obtained by slow evaporation of an ethyl acetate solutio at 283 K.

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Crystal data

 $\begin{array}{l} C_{20}H_{32}N_6O_6\\ M_r = 452.52\\ \text{Monoclinic, } P2_1/c\\ a = 14.6448 \ (15) \ \text{\AA}\\ b = 12.4388 \ (12) \ \text{\AA}\\ c = 13.2805 \ (14) \ \text{\AA}\\ \beta = 98.362 \ (2)^\circ\\ V = 2393.5 \ (4) \ \text{\AA}^3\\ Z = 4 \end{array}$

Data collection

Bruker SMART CCD area-detector
diffractometer
φ and ω scans
Absorption correction: none
11778 measured reflections
4210 independent reflections

Refinement

Refinement on F^2	$w = 1/[\sigma^2(F_0^2) + (0.0815P)^2]$
$R[F^2 > 2\sigma(F^2)] = 0.049$	+ 0.5234P]
$wR(F^2) = 0.148$	where $P = (F_0^2 + 2F_c^2)/3$
S = 1.06	$(\Delta/\sigma)_{\rm max} = 0.001$
4210 reflections	$\Delta \rho_{\rm max} = 0.28 \text{ e } \text{\AA}^{-3}$
297 parameters	$\Delta \rho_{\rm min} = -0.20 \text{ e } \text{\AA}^{-3}$
H-atom parameters constrained	

 $D_x = 1.256 \text{ Mg m}^{-3}$

Cell parameters from 5173

Mo $K\alpha$ radiation

reflections $\theta = 2.3-27.8^{\circ}$

 $\mu = 0.09 \text{ mm}^{-1}$

T = 292 (2) K

 $\begin{aligned} R_{\rm int} &= 0.023\\ \theta_{\rm max} &= 25.0^\circ\\ h &= -16 \rightarrow 17\\ k &= -11 \rightarrow 14\\ l &= -15 \rightarrow 15 \end{aligned}$

Block, colourless

 $0.40 \times 0.30 \times 0.20 \ \text{mm}$

3444 reflections with $I > 2\sigma(I)$

Table 1

Selected geometric parameters (Å, °).

C4-N1	1.497 (3)	C8-N3	1.368 (2)
C5-N1	1.443 (3)	C8-N5	1.393 (3)
C5-N2	1.477 (2)	C9-N3	1.442 (2)
C7-N2	1.368 (2)	C9-N2	1.443 (2)
C7-N4	1.375 (3)		
N2-C7-N4	108.57 (15)	N3-C8-N5	108.26 (15)

All H atoms were treated as riding on their parent C atoms. The methyl H atoms were constrained to an ideal geometry, with C–H distances of 0.96 Å and $U_{\rm iso}(\rm H) = 1.5U_{\rm eq}(\rm C)$, whereas methylene H atoms were constrained to C–H = 0.97 Å with $U_{\rm iso}(\rm H) = 1.5U_{\rm eq}(\rm C)$.

Data collection: *SMART* (Bruker, 2000); cell refinement: *SAINT* (Bruker, 2000); data reduction: *SAINT*; program(s) used to solve



Figure 1

The molecular structure of (I), with the atom-labelling scheme. Displacement ellipsoids are drawn at the 30% probability level and H atoms are represented as spheres of arbitrary radii.

structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *SHELXTL* (Bruker, 2000); software used to prepare material for publication: *SHELXTL*.

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