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#### **Key indicators**

Single-crystal X-ray study T = 292 KMean  $\sigma(\text{C}-\text{C}) = 0.007 \text{ Å}$ Disorder in main residue R factor = 0.072 wR factor = 0.204 Data-to-parameter ratio = 13.1

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

# Diethyl 2,6-dibenzyl-4,8-dioxotetrahydro-2,3a,4a,6,7a,8a-hexaazacyclopenta[*def*]fluorene-8b,8c-dicarboxylate

The molecule of the title compound,  $C_{28}H_{32}N_6O_6$ , shows normal geometrical parameters. The crystal packing is stabilized by a  $C-H\cdots O$  interaction. Received 10 June 2005 Accepted 13 June 2005 Online 17 June 2005

## Comment

Glycoluril derivatives have many areas of applications, such as explosives, slow-release fertilizers, cross-linkers, iodogens, stabilizers of organic compounds against photodegradation, and reagents in combinatorial chemistry (Wu *et al.*, 2002). In this paper, we present the structure of the title compound, (I) (Fig. 1), as a continuation of our previous studies in this area (Wei & Wu, 2005).



All the geometrical parameters for (I) are normal. The crystal packing is stabilized by a C-H···O interaction (Table 1 and Fig. 2), leading to chains of (I) along [100]. The shortest distance between ring centroids of 4.38 Å indicates that any  $\pi$ - $\pi$  effects are very weak.



#### Figure 1

View of (I) showing 30% displacement ellipsoids (arbitrary spheres for the H atoms). Only the major component of the disordered C14/C15 ethyl group is shown.

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# Experimental

Benzylamine (1.07 g, 10 mmol) and formaldehyde (2.4 g, 40 mmol) were added to a stirred solution of diethyl 2,5-dioxotetrahydroimidazo[4,5-*d*]imidazole-3a,6a-dicarboxylate (1.43 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) (2.46 g, 90%) as a colorless solid. Colorless block-like crystals of (I) suitable for data collection were obtained by slow evaporation of a solution in ethyl acetate at 283 K.

#### Crystal data

C <sub>28</sub> H <sub>32</sub> N <sub>6</sub> O <sub>6</sub>	Z = 2
$M_r = 548.60$	$D_x = 1.339 \text{ Mg m}^{-3}$
Triclinic, P1	Mo $K\alpha$ radiation
a = 7.6177 (6) Å	Cell parameters from 4320
b = 12.2882 (9) Å	reflections
c = 14.9197 (11) Å	$\theta = 2.7 - 26.4^{\circ}$
$\alpha = 95.008 \ (1)^{\circ}$	$\mu = 0.10 \text{ mm}^{-1}$
$\beta = 101.415 (1)^{\circ}$	T = 292 (2) K
$\gamma = 92.755 \ (1)^{\circ}$	Block, colorless
V = 1360.69 (18) Å <sup>3</sup>	$0.30 \times 0.20 \times 0.20 \mbox{ mm}$
Data collection	
Bruker SMART CCD diffract-	4689 independent reflections
ometer	3734 reflections with $I > 2\sigma(I)$
$\omega$ and $\varphi$ scans	$R_{\rm int} = 0.021$
Absorption correction: multi-scan	$\theta_{\rm max} = 25.0^{\circ}$
(SADABS; Sheldrick, 1996)	$h = -8 \rightarrow 9$
$T_{\min} = 0.972, T_{\max} = 0.981$	$k = -14 \rightarrow 13$
9666 measured reflections	$l = -17 \rightarrow 17$

## Refinement

Refinement on $F^2$	$w = 1/[\sigma^2(F_o^2) + (0.1005P)^2]$
$R[F^2 > 2\sigma(F^2)] = 0.072$	+ 1.0611P]
$wR(F^2) = 0.204$	where $P = (F_0^2 + 2F_c^2)/3$
S = 1.04	$(\Delta/\sigma)_{\rm max} = 0.002$
4689 reflections	$\Delta \rho_{\rm max} = 0.76 \ {\rm e} \ {\rm \AA}^{-3}$
359 parameters	$\Delta \rho_{\rm min} = -0.42 \text{ e } \text{\AA}^{-3}$
H-atom parameters constrained	

## Table 1

Hydrogen-bond geometry (Å, °).

$D - H \cdot \cdot \cdot A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdots A$
$C8-H8A\cdotsO1^{i}$	0.97	2.44	3.261 (3)	142
6	1.1			

Symmetry codes: (i) x + 1, y, z.

The C14/C15 ethyl group is disordered over two positions in a 0.657 (6):0.343 (6) ratio. All H atoms were positioned geometrically





(C-H = 0.93-0.97 Å) and refined as riding, allowing for free rotation of methyl groups. The constraint  $U_{iso}(H) = 1.2U_{eq}(C)$  or  $1.5U_{eq}(methyl C)$  was applied.

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