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Key indicators

Single-crystal X-ray study
 $T = 150\text{ K}$
Mean $\sigma(\text{C}-\text{C}) = 0.002\text{ \AA}$
 R factor = 0.044
 wR factor = 0.145
Data-to-parameter ratio = 15.1For details of how these key indicators were
automatically derived from the article, see
<http://journals.iucr.org/e>.2,8,12,18-Tetrabutyl-3,7,13,17-tetramethyl-
5,15-bis(3-nitrophenyl)porphyrinogen
methanol disolvate

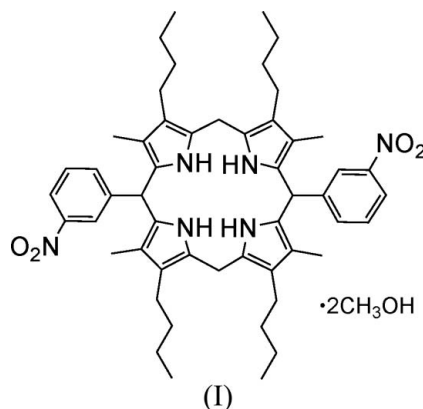
The title compound, $\text{C}_{52}\text{H}_{66}\text{N}_6\text{O}_4 \cdot 2\text{CH}_3\text{OH}$, is a 5,15-diaryl-octaalkylporphyrinogen. The molecule is centrosymmetric, with the pyrrole rings adopting a 1,2-alternate arrangement. The methanol molecules are included in the tetrapyrrole cavity through hydrogen bonding and $\text{O}-\text{H} \cdots \pi$ interactions. The molecules form chains in the crystal structure *via* phenyl $\text{C}-\text{H} \cdots \pi$ -pyrrole stacking.

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Comment

The synthesis of 5,15-diaryloctaalkylporphyrins from the corresponding dipyrromethane is a two-step procedure involving an acid-catalysed reaction with an arylaldehyde to form a porphyrinogen, followed by oxidation of the tetrapyrrolic macrocycle to the porphyrin (Young *et al.*, 1985). During the synthesis of 5,15-bis(3-nitrophenyl)-2,8,12,18-tetrabutyl-3,7,13,17-tetramethylporphyrin from 2,8-dimethyl-3,7-dibutyl-5,10-dihydrodipyrromethane and 3-nitrobenzaldehyde, yellow crystals of the title intermediate porphyrinogen, (I), were formed. The structure of (I) is the first reported for this type of porphyrinogen intermediate bridged by methylene $-\text{CH}_2-$ groups.



The molecular structure of (I) is shown in Fig. 1. The molecule possesses a crystallographic inversion centre. Bond lengths and angles in the porphyrinogen ring core are comparable with those found in similar calix[4]pyrrole derivatives (Gale *et al.*, 1998; Turner *et al.*, 1998). The porphyrinogen ring adopts a 1,2-alternate conformation, similar to that found in calix[4]arenes (Arduini *et al.*, 1997) and some calix[4]pyrroles (Turner *et al.*, 1998). The centrosymmetrically related methanol molecules lie above and below the porphyrinogen ring. These are connected to the macrocycle *via* two hydrogen bonds between methanol atom O3 and the pyrrole H atoms on atoms N1 and N2 (Table 1). Each methanol H atom is centred over the N1-pyrrole ring *via* an

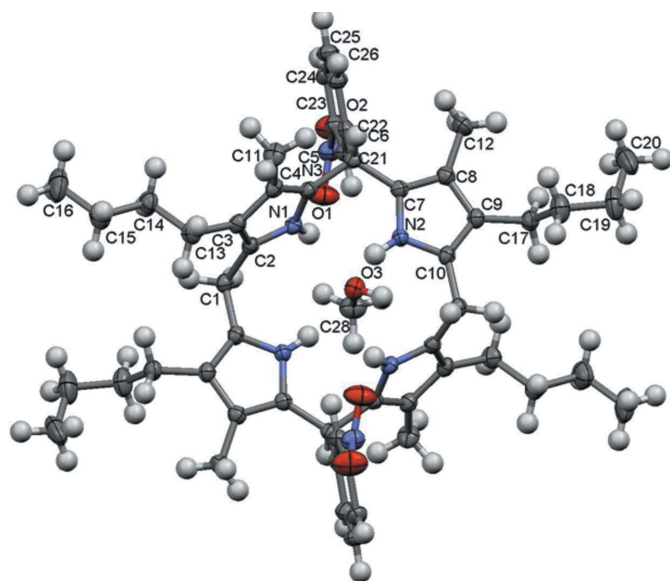


Figure 1
The molecular structure of (I), with 50% probability displacement ellipsoids (arbitrary spheres for H atoms). Unlabelled atoms are related to labelled atoms by the symmetry operator $(-x, -y, -z)$.

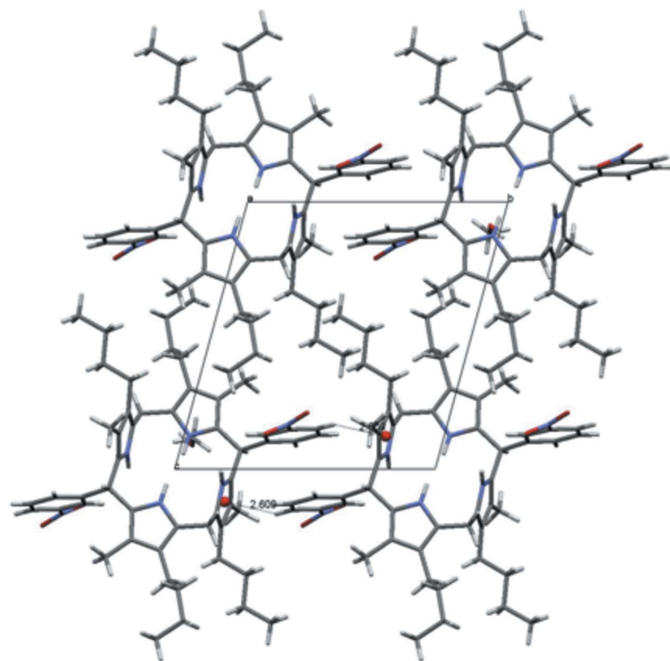


Figure 2
The porphyrin chains of (I), aligned in a two-dimensional assembly, viewed along the a axis. Phenyl $C-H \cdots \pi$ interactions are shown as thin lines and red spheres denote ring centroids.

$O-H \cdots \pi$ interaction, with an $H3 \cdots Cg$ separation of 2.62 Å and an $O3-H3 \cdots Cg$ angle of 131° (Cg is the centroid of the N1-pyrrole ring).

A $C-H \cdots \pi$ interaction between the nitrophenyl group and the pyrrole ring occurs in the crystal structure, with a $C24-H24 \cdots Cg^a$ angle of 150° and an $H24 \cdots Cg^a$ [symmetry code: (a) $-x, -y, -z$] separation of 2.61 Å. Weak intermolecular $C11-H11C \cdots O2$ hydrogen bonding is also observed in the crystal structure (Table 1).

Experimental

A mixture of 4,4'-di-butyl-3,3'-dimethyl-2,2'-dipyrrylmethane (1.04 g, 3.63 mmol) and 3-nitrobenzaldehyde (0.552, 3.65 mmol) in methanol (100 ml) was flushed with nitrogen in the absence of light and cooled in an ice bath for 30 min. To this mixture, trifluoroacetic acid (140 μ l) was added and the solution was stirred for 2 h, during which time yellow crystals of (I) were formed (0.986 g, 32%).

Crystal data

$C_{52}H_{66}N_6O_4 \cdot 2CH_4O$
 $M_r = 903.19$
Triclinic, $P\bar{1}$
 $a = 8.2665$ (1) Å
 $b = 12.2138$ (1) Å
 $c = 13.0914$ (2) Å
 $\alpha = 104.754$ (1) $^\circ$
 $\beta = 97.430$ (1) $^\circ$
 $\gamma = 92.499$ (1) $^\circ$

$V = 1263.42$ (3) Å³
 $Z = 1$
 $D_x = 1.187$ Mg m⁻³
Mo $K\alpha$ radiation
 $\mu = 0.08$ mm⁻¹
 $T = 150$ (2) K
Thick plate, yellow
 $0.34 \times 0.32 \times 0.18$ mm

Data collection

Siemens SMART CCD area-detector diffractometer
 ω scans
Absorption correction: none
9845 measured reflections

4571 independent reflections
4193 reflections with $I > 2\sigma(I)$
 $R_{int} = 0.026$
 $\theta_{max} = 25.3^\circ$

Refinement

Refinement on F^2
 $R[F^2 > 2\sigma(F^2)] = 0.044$
 $wR(F^2) = 0.145$
 $S = 0.81$
4571 reflections
302 parameters
H-atom parameters constrained

$w = 1/[\sigma^2(F_o^2) + (0.1122P)^2 + 0.8341P]$
where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{max} = 0.003$
 $\Delta\rho_{max} = 0.25$ e Å⁻³
 $\Delta\rho_{min} = -0.29$ e Å⁻³

Table 1

Hydrogen-bond geometry (Å, $^\circ$).

$D-H \cdots A$	$D-H$	$H \cdots A$	$D \cdots A$	$D-H \cdots A$
$N1-H1A \cdots O3$	0.88	2.22	2.9930 (16)	147
$N2-H2A \cdots O3^i$	0.88	2.25	3.0881 (17)	160
$C11-H11C \cdots O2^{ii}$	0.98	2.48	3.312 (2)	143

Symmetry codes: (i) $-x, -y, -z$; (ii) $-x, -y + 1, -z$.

Methyl H atoms were placed in calculated positions, with $C-H = 0.98$ Å, and torsion angles were refined to fit the electron density; $U_{iso}(H) = 1.5U_{eq}(C)$. Other H atoms were placed in calculated positions, with $N-H = 0.88$ Å and $C-H = 0.95-1.00$ Å, and were refined using a riding model, with $U_{iso}(H) = 1.2U_{eq}(N,C)$.

Data collection: *SMART* (Siemens, 1995); cell refinement: *SAINT* (Siemens, 1995); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *MERCURY* (Macrae *et al.*, 2006); software used to prepare material for publication: *WinGX* (Farrugia, 1999).

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