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Key indicators

Single-crystal X-ray study T = 130 KMean $\sigma(\text{C-C}) = 0.003 \text{ Å}$ R factor = 0.045 wR factor = 0.117 Data-to-parameter ratio = 18.8

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

Dibenzyl 5-(2-chloroethyl)-3,7-diethyl-2,8-dimethyldipyrromethane-1,9-dicarboxylate

The structure of the title compound [systematic name: dibenzyl 3,3'-diethyl-4,4'-dimethyl-2,2'-(3-chloropropane-1,1-diyl)dipyrrole-5,5'-dicarboxylate], $C_{33}H_{37}ClN_2O_4$, was determined by single-crystal X-ray diffraction at 130 K. In the crystal structure, the compound forms hydrogen-bonded dimers. The two pyrrole H atoms are bonded to one carbonyl O atom of a symmetry-related molecule. Additional, weaker, $C-H\cdots O$ interactions further stabilize the dimers. The two pyrrole rings are inclined to one another by 63.57 (7)°.

Comment

Dipyrromethanes are important building blocks for porphyrin synthesis. Their chemical synthesis in the laboratory often involves the so-called 2 + 2 synthesis, where two dipyrromethane units are condensed (Lindsey, 2000). In nature, porphyrins are prepared by attaching pyrrole units to a dipyrromethane co-factor of porphobilinogen deaminase (Louie *et al.*, 1992; Hädener *et al.*, 1999). Dipyrromethane units are structural building blocks of many bilanes (Zhang *et al.*, 1998; Senge *et al.*, 2001) and calixphyrins (Sessler *et al.*, 2003).



The molecular structure of the title compound, (I), is illustrated in Fig. 1 and selected bond distances and angles are given in Table 1. The two pyrrole rings are inclined to one another by $63.57 (7)^{\circ}$. The molecules form hydrogen-bonded dimers in the crystal structure, utilizing both pyrrole H atoms, but only one of the two carbonyl O atoms. As shown in Fig. 2, carbonyl atom O1 forms two classical hydrogen bonds with the NH units of a neighboring dipyrromethane $(N1-H1\cdots O1^{i})$ and N2-H2 $\cdot \cdot \cdot$ O1ⁱ). A search for weaker hydrogen-bonding interactions (Steiner, 1997) revealed that atom O1 is involved in a third weaker hydrogen bond involving the C51H₂ unit. Thus, each dimer is held together by four classical $N-H \cdots O$ hydrogen bonds and has two C-H···O contacts. Additional close contacts are found within each individual dipyrromethane. Ether atom O2 of one benzyl ester group is separated from the neighboring 2-methyl group by only 2.886 (3) Å, while carbonyl atom O3 of the other benzyl ester group has a close contact to C92 [2.692 (3) Å] and is weakly coordinated to the C81-methyl group. The dimers are further

© 2005 International Union of Crystallography Printed in Great Britain – all rights reserved Received 17 January 2005 Accepted 21 January 2005 Online 29 January 2005 connected by another weak C-H···O interaction. Besides the two intramolecular contacts, O3 is also connected intermolecularly to aromatic atom H14 (H···A = 2.45 Å and D-H···A = 157°) to form infinite chains of the dimers (not shown). Full details of the hydrogen bonding are given in Table 2.

Related crystal structures almost exclusively involve α unsubstituted dipyrromethanes (Lin *et al.*, 1996; Bennis & Gallagher, 1998; Gallagher & Moriarty, 1999; Arumugam *et al.*, 2001; Patra *et al.*, 2002).

Experimental

The compound was prepared as described by Lee & Smith (1997); a sample of the material was kindly provided by Kevin M. Smith (LSU, Baton Rouge) and crystallized from CH_2Cl_2/n -hexane. Crystals were handled as described by Hope (1994).

Z = 2

 $D_x = 1.248 \text{ Mg m}^{-3}$

Cell parameters from 40

 $0.50 \times 0.45 \times 0.39 \text{ mm}$

Mo $K\alpha$ radiation

reflections

T = 130 (2) K

 $\begin{aligned} R_{\rm int} &= 0.041\\ \theta_{\rm max} &= 27.5^\circ \end{aligned}$

 $h = -11 \rightarrow 10$

 $\begin{array}{l} k=-17 \rightarrow 17 \\ l=0 \rightarrow 18 \end{array}$

2 standard reflections

every 198 reflections

intensity decay: 1%

Block, colorless

 $\begin{aligned} \theta &= 19\text{--}23^{\circ} \\ \mu &= 0.17 \text{ mm}^{-1} \end{aligned}$

Crystal data

 $\begin{array}{l} C_{33}H_{37}\text{CIN}_2\text{O}_4 \\ M_r = 561.10 \\ \text{Triclinic, } P\overline{1} \\ a = 8.514 \ (4) \ \text{\AA} \\ b = 13.361 \ (6) \ \text{\AA} \\ c = 14.020 \ (5) \ \text{\AA} \\ \alpha = 92.84 \ (3)^{\circ} \\ \beta = 101.53 \ (3)^{\circ} \\ \gamma = 105.96 \ (3)^{\circ} \\ V = 1493.2 \ (11) \ \text{\AA}^3 \end{array}$

Data collection

Siemens R3m/V diffractometer ω scans Absorption correction: refined from ΔF (*XABS2*; Parkin *et al.*, 1995) $T_{\min} = 0.909$, $T_{\max} = 0.939$ 7317 measured reflections 6854 independent reflections 5327 reflections with $I > 2\sigma(I)$

Refinement

Refinement on F^2	$w = 1/[\sigma^2(F_o^2) + (0.048P)^2]$
$R[F^2 > 2\sigma(F^2)] = 0.045$	+ 0.6397P]
$wR(F^2) = 0.117$	where $P = (F_o^2 + 2F_c^2)/3$
S = 1.02	$(\Delta/\sigma)_{\rm max} < 0.001$
6854 reflections	$\Delta \rho_{\rm max} = 0.26 \text{ e } \text{\AA}^{-3}$
365 parameters	$\Delta \rho_{\rm min} = -0.35 \ {\rm e} \ {\rm \AA}^{-3}$
H-atom parameters constrained	

Table 1

Selected geometric parameters (Å, °).

N1-C4	1.362 (2)	N2-C6	1.356 (2)
N1-C1	1.3793 (19)	N2-C9	1.380 (2)
C1-C2	1.387 (2)	C6-C7	1.391 (2)
C2-C3	1.416 (2)	C7-C8	1.415 (2)
C3-C4	1.389 (2)	C8-C9	1.391 (2)
C4-C5	1.508 (2)	C51-C52	1.506(2)
C5-C6	1.506 (2)	C52-Cl1	1.796 (2)
C5-C51	1.538 (2)		
C4-N1-C1 109.47 (1		C6-N2-C9	109.64 (13)
N1-C4-C5-C6	71.21 (18)	C51-C5-C6-N2	62.76 (19)
C3-C4-C5-C6	-105.77(18)	C4-C5-C6-C7	116.31 (17)
N1-C4-C5-C51	-56.19 (19)	C51-C5-C6-C7	-115.74 (18)
C3-C4-C5-C51	126.83 (17)	C5-C51-C52-Cl1	-62.05 (17)



Figure 1

A view of (I). Displacement ellipsoids are drawn at the 50% probability level; and H atoms have been omitted for clarity.



Figure 2

View of the hydrogen-bonded dimers of (I) in the crystal structure. Full dashed lines indicate strong $N-H\cdots O$ hydrogen bonds, thin dashed lines indicate $C-H\cdots O$ bonding interactions (see Table 2 for details).

Table 2	
Hydrogen-bonding geometry ((Å, °).

$D - H \cdots A$	$D-\mathrm{H}$	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - H \cdots A$
$N1 - H1 \cdots O1^i$	0.88	2.10	2.966 (2)	167
$N2-H2\cdots O1^{i}$	0.88	2.03	2.893 (2)	166
$C5-H5\cdots Cl1$	1.00	2.75	3.204 (2)	108
$C14-H14\cdots O3^{ii}$	0.95	2.45	3.340 (3)	157
$C21 - H21A \cdots O2$	0.98	2.51	2.886 (2)	103
$C51 - H51A \cdots O1^{i}$	0.99	2.57	3.489 (3)	154
C81−H81C···O3	0.98	2.47	3.141 (3)	125
C92−H92 <i>B</i> ···O3	0.99	2.31	2.692 (3)	102

Symmetry codes: (i) 2 - x, 2 - y, 1 - z; (ii) 3 - x, 2 - y, 1 - z.

H atoms were placed in geometrically idealized positions and constrained to ride on their parent atoms, with C-H distances in the range 0.95–1.00 Å and N-H distances of 0.88 Å, and with $U_{iso}(H) = 1.2U_{eq}(C,N)$ for CH, CH₂ and pyrrole H atoms, and $U_{iso}(H) = 1.5U_{eq}(C)$ for methyl groups.

Data collection: *P3* (Siemens, 1995); cell refinement: *P3*; data reduction: *XDISK* (Siemens, 1995); program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *XP* (Siemens, 1995); software used to prepare material for publication: *SHELXL97* and *PLATON* (Spek, 2003).

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