# organic papers

Acta Crystallographica Section E Structure Reports Online

ISSN 1600-5368

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#### **Key indicators**

Single-crystal X-ray study T = 293 K Mean  $\sigma$ (C–C) = 0.003 Å R factor = 0.054 wR factor = 0.161 Data-to-parameter ratio = 18.7

For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e. 1,8-Bis(3-formyl-5-methyl-2-oxidobenzyl)-1,8-diaza-4,11-diazoniacyclotetradecane chloroform solvate

## Comment

The special feature of the title ligand, (I), is the presence of a formyl group which can be further condensed with polymers. These macrocycle-appended polymers can be used in metal extraction processes (Bianchi *et al.*, 1991) and catalytic applications. The aldehyde group can also be condensed with a fluorescent unit (Kawakami *et al.*, 2001) or redox-active molecules which can be used as sensors (Yamamoto *et al.*, 1996). As the title compound, (I), is of high industrial value, we have undertaken the crystal structure determination by X-ray diffraction.



The asymmetric unit of (I) contains two independent halfmacrocycles and one solvent (chloroform) molecule. One-half of each macrocycle is inversion-related to the other half (Fig. 1). The C–C and N–C bond lengths agree well with the reported literature values (Allen *et al.*, 1987). The sums of the bond angles around N1 [334.1°] and N3 [335.3°] indicate  $sp^3$ hybridization. The exocyclic angles around C6 and C20 show considerable asymmetry, with the C9–C6–C7 [122.09 (17)°] and C23–C20–C21 [121.16 (16)°] angles larger than the C9–C6–C5 [117.97 (15)°] and C23–C20–C19 [118.33 (15)°] angles.

The conformation of the macrocycle is stabilized by N– H···O and C–H···O intramolecular interactions (Table 1). The molecular structure is also stabilized by intramolecular C–H···O interactions, and the crystal packing is stabilized by C–H···O and C–H··· $\pi$  intermolecular interactions. Received 10 July 2006 Accepted 31 July 2006

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## **Experimental**

3-chloromethyl-5-methylsalicylaldehyde 5-Methylsalicylaldehyde, 1,4,8,11-tetrazatricyclo[9.3.1.1]hexadecane were prepared and according to the methods of Duff (1941), Royal et al. (1998) and Gane et al. (1991), respectively. 1,4,8,11-Tetrazatricyclo[9.3.1.1]tetradecane (1 g, 0.004 mol) was dissolved in acetonitrile (30 ml) and two equivalents of 3-chloromethyl-5-methylsalicylaldehyde (1.64 g, 0.0089 mol) in acetonitrile (30 ml) were rapidly added; the solution was stirred at 298 K for 24 h. The precipitate formed was filtered off, washed with acetonitrile and dried under vacuum. The resulting crude compound, 1,8-N,N'-bis[(3-formyl-2-hydroxy-5-methyl)benzyl]-4,11-diatricyclo[9.3.1.1]tetradecane dichloride, was recrystallized from water to give white crystals (yield 85%, m.p. 558 K). This compound (0.5 g) was dissolved in an aqueous NaOH solution  $(100 \text{ ml}, 3 \times 10^{-1} \text{ M})$  with stirring. After stirring for 4 h, the solution was extracted with  $CHCl_3$  (5 × 50 ml). The combined  $CHCl_3$  extracts were dried with anhydrous MgSO4 and concentrated under vacuum to obtain a light-yellow compound, (I). Light-yellow crystals of the title compound were obtained upon recrystallization from chloroform (yield 70%, m.p. 573 K).

### Crystal data

C28H40N4O4·CHCl3	$V = 1539.74 (12) \text{ Å}^3$
$M_r = 616.01$	Z = 2
Triclinic, $P\overline{1}$	$D_x = 1.329 \text{ Mg m}^{-3}$
a = 11.1366 (5)  Å	Mo $K\alpha$ radiation
b = 11.2816 (5) Å	$\mu = 0.34 \text{ mm}^{-1}$
c = 14.2124 (7) Å	T = 293 (2) K
$\alpha = 69.564 \ (1)^{\circ}$	Block, yellow
$\beta = 84.559 \ (1)^{\circ}$	$0.29 \times 0.26 \times 0.25 \text{ mm}$
$\gamma = 67.110 \ (1)^{\circ}$	

Data collection

Bruker SMART APEX CCD areadetector diffractometer ω scans Absorption correction: none 17829 measured reflections

#### Refinement

Refinement on $F^2$	$w = 1/[\sigma^2(F_0^2) + (0.0823P)^2]$
$R[F^2 > 2\sigma(F^2)] = 0.054$	+ 0.5118P]
$wR(F^2) = 0.161$	where $P = (F_0^2 + 2F_c^2)/3$
S = 1.07	$(\Delta/\sigma)_{\rm max} = 0.001$
7069 reflections	$\Delta \rho_{\rm max} = 0.69 \ {\rm e} \ {\rm \AA}^{-3}$
379 parameters	$\Delta \rho_{\rm min} = -0.43 \text{ e } \text{\AA}^{-3}$
H atoms treated by a mixture of	
independent and constrained	
refinement	

7069 independent reflections

 $R_{\rm int} = 0.018$ 

 $\theta_{\rm max} = 28.0^\circ$ 

5878 reflections with  $I > 2\sigma(I)$ 

## Table 1

Hydrogen-bond geometry (Å, °).

Cg1 is the centroid of the C16-C21 ring.

$D - H \cdots A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - H \cdot \cdot \cdot A$
$N2-H2A\cdots O2$	0.90 (2)	1.71 (2)	2.605 (2)	169 (2)
$N2-H2B\cdots N1^{i}$	0.90 (2)	2.26 (2)	2.909 (2)	128 (2)
$N4-H4A\cdots O4$	0.90 (2)	1.85 (2)	2.741 (2)	171 (2)
$N4-H4B\cdots O4^{ii}$	0.90 (2)	2.01 (2)	2.883 (2)	163 (2)
$N4-H4B\cdots N3^{ii}$	0.90 (2)	2.47 (2)	2.984 (2)	117 (2)
C8-H8···O2	0.93	2.46	2.808 (3)	102
$C11-H11A \cdot \cdot \cdot O2^{i}$	0.97	2.50	3.231 (3)	133
$C14-H14B\cdots O4^{iii}$	0.97	2.59	3.557 (3)	172
C22-H22···O4	0.93	2.49	2.828 (3)	102
$C26-H26A\cdots O4^{ii}$	0.97	2.56	3.213 (2)	125



Figure 1

The structure of (I), showing 30% probability displacement ellipsoids. Atoms labelled with the suffixes a and b are generated by the symmetry operations (1 - x, 1 - y, 1 - z) and (1 - x, 2 - y, 2 - z), respectively. Hydrogen bonds are shown as dashed lines and H atoms not involved in hydrogen bonding have been omitted.

$D - H \cdot \cdot \cdot A$	$D-\mathrm{H}$	$H \cdot \cdot \cdot A$	$D \cdots A$	D-H	$\cdot \cdot \cdot A$
$C27-H27B\cdots O1^{ii}$ $C29-H29\cdots O1^{ii}$ $C13-H13A\cdots Cg1^{iii}$	0.97 0.98 0.97	2.39 2.46 2.58	3.317 (3) 3.326 (3) 3.483 (2)	161 148 154	
Symmetry codes: (i) $x, y - 1, z$ .	-x + 1, -y +	-1, -z + 1; (ii)	) $-x+1, -y+2$	2, -z + 2;	(iii)

The amine H atoms were located in a difference map and refined with an N-H distance restraint of 0.90 (1) Å. The C-bound H atoms were positioned geometrically and allowed to ride on their parent C atoms, with C-H = 0.93-0.97 Å and  $U_{iso}(H) = 1.2U_{eq}(C)$  or  $1.5U_{eq}(C_{methyl})$ .

Data collection: *SMART* (Bruker, 2001); cell refinement: *SAINT* (Bruker, 2001); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *PLATON* (Spek, 2003); software used to prepare material for publication: *SHELXL97* and *PARST* (Nardelli, 1995).

DG and DV acknowledge financial support from the University Grants Commission (UGC-SAP) and the Department of Science and Technology (DST-FIST), Government of India.

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