organic papers

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Key indicators

Single-crystal X-ray study T = 173 K Mean σ (C–C) = 0.004 Å R factor = 0.043 wR factor = 0.130 Data-to-parameter ratio = 20.6

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

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4,11-Bis(acetamido)-7,14-dimethyl-4,11-diaza-1,8diazoniacyclotetradecane diperchlorate dihydrate

The title compound features a doubly protonated dimethyland bis(methylcarboxamide)-substituted tetraazacyclotetradecane with the pendant arms disposed over the cavity, an arrangement that is facilitated by intramolecular hydrogenbonding interactions. The dication occupies a special position about an inversion centre, whereas the ClO_4^- anion and the water molecule are in general positions, thus forming a crystal with the overall composition $C_{16}H_{36}N_6O_2^{2+}\cdot 2\text{ClO}_4^-\cdot 2H_2O$. The crystal structure is stabilized by a variety of hydrogenbonding contacts involving the dication, perchlorate anion and solvent water molecule.

Comment

The crystal structure of the title compound, (I), comprises dications, perchlorate anions and solvent water molecules in the ratio 1:2:2. The cation (Fig. 1) occupies a special position about an inversion centre so that the four N atoms within the macrocycle are coplanar. Protonation occurs at the N1 atom and this is reflected in the systematic differences in the N1–C and N4–C bond distances (Table 1). The pendant acetamide groups lie to either side of the N₄ plane and are orientated so as to place the carbonyl O atoms over the cavity of the macrocycle.



This configuration is stabilized by hydrogen-bonding interactions involving the N1-bound H1n atoms which are directed towards the inside of the cavity. Thus, O4" is equally separated from H1n and the symmetry-related H1nⁱ atoms, indicating that each H1n atom is bifurcated and that a hydrogen-bonded H₂O₂ square is formed perpendicular to the N₄ plane [symmetry code: (i) -x, -y, -z]. Geometric details of the hydrogen-bonding system in the structure of the title compound are given in Table 2.

The O4" atom also forms an acceptor interaction with the solvent water molecule and the other water-bound H atom

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 $h = -21 \rightarrow 8$ $k = 0 \rightarrow 14$ $l = -18 \rightarrow 18$

3 standard reflections

every 400 reflections

intensity decay: 1.8%

H-atom parameters constrained

 $w = 1/[\sigma^2(F_o^2) + (0.061P)^2]$

 $(\Delta/\sigma)_{\rm max} < 0.001$ $\Delta \rho_{\rm max} = 0.30 \text{ e} \text{ Å}^{-3}$

 $\Delta \rho_{\rm min} = -0.38 \text{ e} \text{ Å}^{-3}$

where $P = (F_o^2 + 2F_c^2)/3$





The molecular structure and crystallographic numbering scheme for the dication in (I). Displacement ellipsoids are shown at the 50% probability level (Johnson, 1976).

forms a donor interaction to a perchlorate O atom. The solvent water molecule is also involved in an acceptor interaction with the cationic NH2 group of the macrocycle. Each of the amide H atoms forms a donor interaction with one of the perchlorate O atoms.

The aforementioned hydrogen-bonding network results in a structure that may be described as being comprised of alternating stacks of cations, and anions and water molecules aligned along the *c* direction.

Experimental

The 1,8-bis(acetamido)-7,14-dimethyl-1,4,8,11-tetraazacyclotetradecane compound was prepared in the following manner: 7,14-dimethyl-1,4,8,11-tetraazacyclotetradecane (Fairbank et al., 1985) (2.28 g, 0.01 mol) in ethanol (20 ml) was added to a solution of 2-chloroacetamide (4.68 g, 0.05 mol) and triethylamine (6.07 g, 0.06 mol) also in ethanol (30 ml). The mixture was then refluxed for one day. After the solution had been allowed to stand for three days, a quantity of colourless crystals precipitated. These were recrystallized from a water/acetonitrile (1:1, 20 ml) mixture. Crystals of the title compound were isolated as colourless blocks from the attempted reaction of 1,8bis(acetamido)-7,14-dimethyl-1,4,8,11-tetraazacyclotetradecane and zinc perchlorate in aqueous medium.

Crystal data

$C_{16}H_{36}N_6O_2^{2+}\cdot 2ClO_4^{-}\cdot 2H_2O$
$M_r = 579.43$
Monoclinic, C2/c
a = 16.576 (3) Å
b = 10.877 (6) Å
c = 14.314 (2) Å
$\beta = 90.25 \ (1)^{\circ}$
$V = 2581 (1) \text{ Å}^3$
$\mathbf{Z} = \mathbf{A}$

 $D_x = 1.491 \text{ Mg m}^{-3}$ Mo $K\alpha$ radiation Cell parameters from 20 reflections $\theta = 7.5 - 14.6^{\circ}$ $\mu = 0.32 \text{ mm}^{-1}$ T = 173 KBlock, colourless $0.36 \times 0.23 \times 0.13 \mbox{ mm}$

Data collection	
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1 ł

2

Rigaku AFC-7R diffractometer
ω –2 θ scans
4647 measured reflections
2965 independent reflections
1545 reflections with $I > 2\sigma(I)$
$R_{\rm int} = 0.049$
$\theta_{\rm max} = 27.5^{\circ}$

Refinement

Refinement on F^2 $R[F^2 > 2\sigma(F^2)] = 0.043$ $wR(F^2) = 0.130$ S = 0.982965 reflections 144 parameters

Table 1

Selected geometric parameters (Å, °).

O4''-C4''	1.241 (3)	N4''-C4''	1.331 (4)
N1-C2	1.495 (4)	C2-C3	1.519 (4)
N1-C7 ⁱ	1.507 (3)	C4′-C4′′	1.516 (4)
N4-C3	1.471 (3)	C5-C6	1.521 (4)
N4-C4′	1.463 (3)	C6-C7	1.521 (4)
N4-C5	1.481 (3)	C7-C7′	1.520 (4)
C2-N1-C7 ⁱ	114.7 (2)	O4''-C4''-C4'	120.9 (2)
C4'-N4-C3	111.7 (2)	N4''-C4''-C4'	115.9 (3)
C4'-N4-C5	110.2 (2)	N4-C5-C6	113.3 (2)
C3-N4-C5	108.6 (2)	C7-C6-C5	114.8 (2)
N1-C2-C3	111.7 (2)	C7′-C7-C6	112.2 (2)
N4-C3-C2	114.0 (2)	C7'-C7-N1 ⁱ	111.1 (2)
N4-C4'-C4''	113.3 (2)	C6-C7-N1 ⁱ	108.1 (2)
O4''-C4''-N4''	123.2 (3)		

Symmetry code: (i) -x, -y, -z.

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

$D - H \cdot \cdot \cdot A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdots A$
$N1-H1n\cdots O4''$	0.93	2.53	3.083 (3)	118
$N1-H1n\cdots O4''^i$	0.93	2.54	3.079 (3)	118
$O1w - H2o \cdots O4''$	0.99	1.76	2.744 (3)	176
$O1w - H1o \cdots O12^{ii}$	0.90	2.02	2.903 (3)	166
$N1 - H2n \cdot \cdot \cdot O1w^{iii}$	0.88	1.92	2.768 (3)	163
$N4'' - H3n \cdot \cdot \cdot O13^{iv}$	0.95	2.27	3.110 (4)	148
$N4'' - H4n \cdots O11^v$	0.95	2.26	3.122 (4)	151
Summetry codes: (i)		(ii) $-\frac{1}{2} - x \frac{1}{2}$	- v 7: (iii) x	$-v^{\frac{1}{2}} + z$; (iv)

Symmetry codes: (1) -x, -y, -z; (ii) $-\frac{1}{2} - x, \frac{1}{2} - y, -z$; (iii) $\overline{x, -y, \frac{1}{2} + z}$; (iv) $x, 1 - y, z - \frac{1}{2}$; (v) -x, 1 - y, -z.

The C-bound H atoms were placed in geometrically calculated positions and included in the final refinement in the riding-model approximation with an overall displacement parameter. The O- and N-bound H atoms were located from a difference Fourier map but were not refined.

Data collection: MSC/AFC Diffractometer Control Software (Molecular Structure Corporation, 1996); cell refinement: MSC/AFC Diffractometer Control Software; data reduction: TEXSAN for Windows (Molecular Structure Corporation, 1997); program(s) used to solve structure: PATTY in DIRDIF92 (Beurskens et al., 1992); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); software used to prepare material for publication: SHELXL97.

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