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Octane-1,8-diylpyridinium dibromide dihydrate

 Xin Xiao,^a Ping-Yue Jiang,^a Yun-Qian Zhang,^{a*} Sai-Feng Xue^b and Qian-Jiang Zhu^b
^aKey Laboratory of Macrocyclic and Supramolecular Chemistry of Guizhou Province, Guizhou University, Guiyang 550025, People's Republic of China, and ^bInstitute of Applied Chemistry, Guizhou University, Guiyang 550025, People's Republic of China

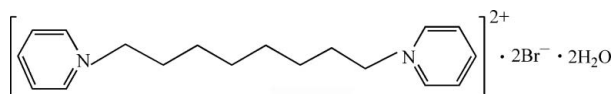
Correspondence e-mail: sci.yqzhang@gzu.edu.cn

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 Key indicators: single-crystal X-ray study; $T = 273$ K; mean $\sigma(\text{C}-\text{C}) = 0.003$ Å; R factor = 0.029; wR factor = 0.077; data-to-parameter ratio = 23.0.

The asymmetric unit of the title compound, $\text{C}_{18}\text{H}_{26}\text{N}_2^{2+} \cdot 2\text{Br}^- \cdot 2\text{H}_2\text{O}$, consists of one-half of the organic cation, one Br^- anion and one water molecule. The organic cation is situated on a centre of inversion. The dihedral angle between the pyridine ring and the plane of the central linkage is $59.3(1)^\circ$. The cations, anions and water molecules are linked *via* $\text{O}-\text{H} \cdots \text{Br}$, $\text{C}-\text{H} \cdots \text{Br}$ and $\text{C}-\text{H} \cdots \text{O}$ hydrogen bonds, forming a three-dimensional framework.

Related literature

 For general background, see: Day *et al.* (2000, 2002); Freeman *et al.* (1981); Kim *et al.* (2000).


Experimental

Crystal data

 $\text{C}_{18}\text{H}_{26}\text{N}_2^{2+} \cdot 2\text{Br}^- \cdot 2\text{H}_2\text{O}$
 $M_r = 466.26$

 Orthorhombic, *Pbca*
 $a = 9.8329(7)$ Å

 $b = 13.3000(11)$ Å

 $c = 16.5688(13)$ Å

 $V = 2166.8(3)$ Å³
 $Z = 4$

 Mo $K\alpha$ radiation

 $\mu = 3.75$ mm⁻¹
 $T = 273(2)$ K

 $0.32 \times 0.24 \times 0.19$ mm

Data collection

Bruker APEXII CCD area-detector diffractometer

 Absorption correction: multi-scan (*SADABS*; Bruker, 2005)

 $T_{\min} = 0.380$, $T_{\max} = 0.536$

(expected range = 0.347–0.490)

21410 measured reflections

2503 independent reflections

 1842 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.034$

Refinement

 $R[F^2 > 2\sigma(F^2)] = 0.029$
 $wR(F^2) = 0.077$
 $S = 1.01$

2503 reflections

109 parameters

H-atom parameters constrained

 $\Delta\rho_{\text{max}} = 0.25$ e Å⁻³
 $\Delta\rho_{\text{min}} = -0.39$ e Å⁻³

Table 1

Hydrogen-bond geometry (Å, °).

$D-H \cdots A$	$D-H$	$H \cdots A$	$D \cdots A$	$D-H \cdots A$
$\text{O1W}-\text{H1WB} \cdots \text{Br1}$	0.81	2.55	3.353 (2)	173
$\text{O1W}-\text{H1WA} \cdots \text{Br1}^i$	0.84	2.57	3.392 (2)	169
$\text{C1}-\text{H1} \cdots \text{Br1}$	0.93	2.82	3.596 (2)	141
$\text{C2}-\text{H2} \cdots \text{O1W}^{ii}$	0.93	2.48	3.271 (3)	143
$\text{C5}-\text{H5} \cdots \text{Br1}^{iii}$	0.93	2.67	3.588 (2)	168

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

Data collection: *APEX2* (Bruker, 2005); cell refinement: *SAINT* (Bruker, 2005); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *ORTEP-3 for Windows* (Farrugia, 1997); software used to prepare material for publication: *WinGX* (Farrugia, 1999).

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: CI2531).

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supporting information

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Octane-1,8-diylidipyridinium dibromide dihydrate

Xin Xiao, Ping-Yue Jiang, Yun-Qian Zhang, Sai-Feng Xue and Qian-Jiang Zhu

S1. Comment

As part of our ongoing investigation on bipyridyl compounds, we present here the crystal structure of the title compound, (I), which can develop strong intermolecular interactions with cucurbit[*n*]urils (CB[*n*]) (Freeman *et al.*, 1981; Day *et al.*, 2000, 2002; Kim *et al.*, 2000).

The asymmetric unit of compound (I) (Fig. 1) consists of one-half of the organic cation, one Br⁻ anion and one lattice water molecule. The organic cation is situated on a centre of inversion which coincides with the midpoint of the C9—C9ⁱ bond [symmetry code: (i) 1 - x, 1 - y, -z]. The two pyridine rings are parallel by virtue of the centre of symmetry. The dihedral angle between the pyridine ring and the central C6—C9/C6ⁱ—C9ⁱ chain is 59.3 (1)°. The cations, anions and water molecules are linked *via* O—H...Br, C—H...Br and C—H...O hydrogen bonds (Table 1) forming a three-dimensional framework.

S2. Experimental

A solution of 1,8-dibromine-octane (2.72 g, 0.01 mol) was added to a stirred solution of pyridine (1.98 g, 0.025 mol) in 1,4-dioxane (50 ml) at 383 K for 5 h. After cooling to room temperature, the mixture was filtered. The solid product was dissolved in 80 ml water, and then set aside for four weeks to obtain colourless crystals of the title compound.

S3. Refinement

Water H atoms were located in a difference Fourier map and refined as riding in their as-found positions relative to the parent O atom, with $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{O})$. All other H atoms were placed in calculated positions and refined as riding, with C—H = 0.93–0.97 Å and $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$.

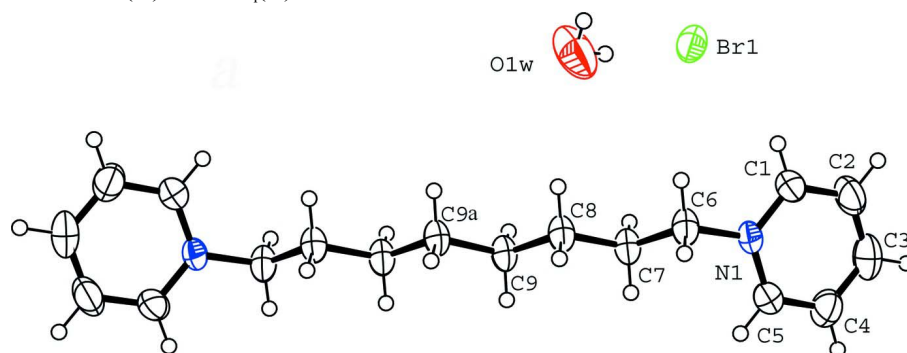


Figure 1

The molecular structure of (I), showing the atom-labelling scheme for the asymmetric unit. Displacement ellipsoids are drawn at the 50% probability level. unlabelled atoms are related to labelled atoms by the symmetry operation (1 - x, 1 - y, -z). Symmetry related bromide ion and water molecule are not shown.

Octane-1,8-diylidpyridinium dibromide dihydrate

Crystal data

 $C_{18}H_{26}N_2^{2+} \cdot 2(Br^-) \cdot 2H_2O$ $M_r = 466.26$ Orthorhombic, *Pbca*

Hall symbol: -P 2ac 2ab

 $a = 9.8329$ (7) Å $b = 13.3000$ (11) Å $c = 16.5688$ (13) Å $V = 2166.8$ (3) Å³ $Z = 4$ $F(000) = 952$ $D_x = 1.429$ Mg m⁻³Mo $K\alpha$ radiation, $\lambda = 0.71073$ Å

Cell parameters from 21410 reflections

 $\theta = 2.5$ – 27.6° $\mu = 3.75$ mm⁻¹ $T = 273$ K

Prism, colourless

 $0.32 \times 0.24 \times 0.19$ mm

Data collection

Bruker APEXII CCD area-detector
diffractometer

Radiation source: fine-focus sealed tube

Graphite monochromator

 φ and ω scansAbsorption correction: multi-scan
(*SADABS*; Bruker, 2005) $T_{\min} = 0.380$, $T_{\max} = 0.536$

21410 measured reflections

2503 independent reflections

1842 reflections with $I > 2\sigma(I)$ $R_{\text{int}} = 0.034$ $\theta_{\max} = 27.6^\circ$, $\theta_{\min} = 2.5^\circ$ $h = -12 \rightarrow 12$ $k = -17 \rightarrow 17$ $l = -15 \rightarrow 21$

Refinement

Refinement on F^2

Least-squares matrix: full

 $R[F^2 > 2\sigma(F^2)] = 0.029$ $wR(F^2) = 0.077$ $S = 1.01$

2503 reflections

109 parameters

0 restraints

Primary atom site location: structure-invariant
direct methodsSecondary atom site location: difference Fourier
mapHydrogen site location: inferred from
neighbouring sites

H-atom parameters constrained

 $w = 1/[\sigma^2(F_o^2) + (0.0352P)^2 + 0.8723P]$ where $P = (F_o^2 + 2F_c^2)/3$ $(\Delta/\sigma)_{\max} = 0.001$ $\Delta\rho_{\max} = 0.25$ e Å⁻³ $\Delta\rho_{\min} = -0.39$ e Å⁻³

Special details

Geometry. All e.s.d.'s (except the e.s.d. in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell e.s.d.'s are taken into account individually in the estimation of e.s.d.'s in distances, angles and torsion angles; correlations between e.s.d.'s in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell e.s.d.'s is used for estimating e.s.d.'s involving l.s. planes.

Refinement. Refinement of F^2 against ALL reflections. The weighted R -factor wR and goodness of fit S are based on F^2 , conventional R -factors R are based on F , with F set to zero for negative F^2 . The threshold expression of $F^2 > \sigma(F^2)$ is used only for calculating R -factors(gt) *etc.* and is not relevant to the choice of reflections for refinement. R -factors based on F^2 are statistically about twice as large as those based on F , and R -factors based on ALL data will be even larger.

Fractional atomic coordinates and isotropic or equivalent isotropic displacement parameters (Å²)

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$
Br1	0.28891 (2)	0.652327 (19)	0.328729 (15)	0.05280 (11)
N1	0.44180 (18)	0.35645 (13)	0.32104 (10)	0.0376 (4)
C1	0.3916 (3)	0.40288 (17)	0.38614 (14)	0.0507 (6)
H1	0.3246	0.4517	0.3801	0.061*
C2	0.4380 (3)	0.3791 (2)	0.46131 (16)	0.0642 (8)

H2	0.4014	0.4105	0.5065	0.077*
C3	0.5389 (3)	0.3088 (2)	0.46995 (16)	0.0660 (8)
H3	0.5725	0.2931	0.5209	0.079*
C4	0.5895 (3)	0.2622 (2)	0.40288 (15)	0.0588 (7)
H4	0.6582	0.2146	0.4077	0.071*
C5	0.5382 (2)	0.28617 (18)	0.32886 (13)	0.0475 (6)
H5	0.5707	0.2532	0.2832	0.057*
C6	0.3864 (2)	0.37965 (18)	0.23978 (13)	0.0455 (5)
H6A	0.3137	0.4285	0.2453	0.055*
H6B	0.3476	0.3189	0.2170	0.055*
C7	0.4916 (2)	0.42052 (18)	0.18213 (12)	0.0432 (5)
H7A	0.5633	0.3712	0.1749	0.052*
H7B	0.5320	0.4808	0.2048	0.052*
C8	0.4286 (2)	0.44515 (19)	0.10078 (13)	0.0444 (5)
H8A	0.3865	0.3850	0.0791	0.053*
H8B	0.3577	0.4950	0.1084	0.053*
C9	0.5310 (2)	0.48488 (17)	0.04014 (12)	0.0408 (5)
H9A	0.5766	0.5428	0.0633	0.049*
H9B	0.5992	0.4335	0.0306	0.049*
O1W	0.1270 (2)	0.6061 (2)	0.15455 (12)	0.0973 (8)
H1WA	0.0442	0.6131	0.1647	0.117*
H1WB	0.1689	0.6117	0.1965	0.117*

Atomic displacement parameters (Å²)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
Br1	0.04958 (16)	0.05876 (17)	0.05006 (17)	0.00012 (12)	-0.00083 (11)	0.01642 (11)
N1	0.0424 (10)	0.0428 (10)	0.0277 (9)	-0.0036 (8)	0.0058 (7)	0.0044 (7)
C1	0.0627 (14)	0.0464 (13)	0.0429 (14)	0.0058 (12)	0.0161 (12)	0.0025 (11)
C2	0.094 (2)	0.0637 (16)	0.0345 (14)	-0.0017 (16)	0.0104 (14)	-0.0069 (12)
C3	0.081 (2)	0.0830 (19)	0.0341 (14)	-0.0090 (17)	-0.0084 (14)	0.0108 (14)
C4	0.0601 (15)	0.0672 (17)	0.0492 (15)	0.0092 (13)	-0.0038 (13)	0.0141 (13)
C5	0.0528 (14)	0.0527 (14)	0.0368 (13)	0.0072 (11)	0.0061 (11)	0.0008 (11)
C6	0.0437 (12)	0.0601 (13)	0.0326 (12)	-0.0035 (11)	-0.0004 (10)	0.0098 (11)
C7	0.0447 (12)	0.0539 (13)	0.0311 (12)	-0.0047 (11)	0.0010 (10)	0.0064 (10)
C8	0.0437 (12)	0.0590 (14)	0.0305 (11)	-0.0035 (10)	-0.0010 (10)	0.0069 (10)
C9	0.0402 (11)	0.0540 (13)	0.0281 (11)	-0.0067 (10)	-0.0031 (9)	0.0036 (10)
O1W	0.0625 (12)	0.177 (3)	0.0519 (12)	0.0086 (16)	-0.0006 (10)	-0.0161 (14)

Geometric parameters (Å, °)

N1—C1	1.337 (3)	C6—H6A	0.97
N1—C5	1.337 (3)	C6—H6B	0.97
N1—C6	1.485 (3)	C7—C8	1.520 (3)
C1—C2	1.364 (4)	C7—H7A	0.97
C1—H1	0.93	C7—H7B	0.97
C2—C3	1.370 (4)	C8—C9	1.518 (3)
C2—H2	0.93	C8—H8A	0.97

C3—C4	1.366 (4)	C8—H8B	0.97
C3—H3	0.93	C9—C9 ⁱ	1.518 (4)
C4—C5	1.364 (3)	C9—H9A	0.97
C4—H4	0.93	C9—H9B	0.97
C5—H5	0.93	O1W—H1WA	0.84
C6—C7	1.509 (3)	O1W—H1WB	0.81
C1—N1—C5	120.42 (19)	C7—C6—H6B	108.9
C1—N1—C6	120.0 (2)	H6A—C6—H6B	107.7
C5—N1—C6	119.53 (18)	C6—C7—C8	111.07 (19)
N1—C1—C2	120.4 (2)	C6—C7—H7A	109.4
N1—C1—H1	119.8	C8—C7—H7A	109.4
C2—C1—H1	119.8	C6—C7—H7B	109.4
C1—C2—C3	119.7 (2)	C8—C7—H7B	109.4
C1—C2—H2	120.1	H7A—C7—H7B	108.0
C3—C2—H2	120.1	C9—C8—C7	113.04 (18)
C4—C3—C2	119.2 (2)	C9—C8—H8A	109.0
C4—C3—H3	120.4	C7—C8—H8A	109.0
C2—C3—H3	120.4	C9—C8—H8B	109.0
C5—C4—C3	119.4 (2)	C7—C8—H8B	109.0
C5—C4—H4	120.3	H8A—C8—H8B	107.8
C3—C4—H4	120.3	C9 ⁱ —C9—C8	113.9 (2)
N1—C5—C4	120.8 (2)	C9 ⁱ —C9—H9A	108.8
N1—C5—H5	119.6	C8—C9—H9A	108.8
C4—C5—H5	119.6	C9 ⁱ —C9—H9B	108.8
N1—C6—C7	113.42 (17)	C8—C9—H9B	108.8
N1—C6—H6A	108.9	H9A—C9—H9B	107.7
C7—C6—H6A	108.9	H1WA—O1W—H1WB	108.2
N1—C6—H6B	108.9		
C5—N1—C1—C2	0.0 (3)	C3—C4—C5—N1	1.8 (4)
C6—N1—C1—C2	177.4 (2)	C1—N1—C6—C7	120.1 (2)
N1—C1—C2—C3	1.5 (4)	C5—N1—C6—C7	-62.5 (3)
C1—C2—C3—C4	-1.3 (4)	N1—C6—C7—C8	-178.7 (2)
C2—C3—C4—C5	-0.3 (4)	C6—C7—C8—C9	-179.0 (2)
C1—N1—C5—C4	-1.6 (3)	C7—C8—C9—C9 ⁱ	-177.0 (2)
C6—N1—C5—C4	-179.0 (2)		

Symmetry code: (i) $-x+1, -y+1, -z$.

Hydrogen-bond geometry (\AA , $^\circ$)

<i>D</i> —H \cdots <i>A</i>	<i>D</i> —H	H \cdots <i>A</i>	<i>D</i> \cdots <i>A</i>	<i>D</i> —H \cdots <i>A</i>
O1W—H1WB \cdots Br1	0.81	2.55	3.353 (2)	173
O1W—H1WA \cdots Br1 ⁱⁱ	0.84	2.57	3.392 (2)	169
C1—H1 \cdots Br1	0.93	2.82	3.596 (2)	141

C2—H2...O1 ⁱⁱ	0.93	2.48	3.271 (3)	143
C5—H5...Br1 ^{iv}	0.93	2.67	3.588 (2)	168

Symmetry codes: (ii) $x-1/2, y, -z+1/2$; (iii) $-x+1/2, -y+1, z+1/2$; (iv) $-x+1, y-1/2, -z+1/2$.