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Key indicators

Single-crystal X-ray study T = 170 K Mean σ (C–C) = 0.003 Å R factor = 0.036 wR factor = 0.101 Data-to-parameter ratio = 13.7

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

2,2'-(*p*-Phenylenediimino)dipyridinium dichloride hexahydrate

In the crystal structure of the title compound, $C_{16}H_{16}N_4^{2+} \cdot 2C I^- \cdot 6H_2O$, the 2,2'-(*p*-phenylenediimino)dipyridinium cation has inversion symmetry. The chloride ions and water molecules are linked *via* $O-H \cdot \cdot \cdot O$ and $O-H \cdot \cdot \cdot CI^-$ hydrogen bonds into an infinite two-dimensional network parallel to the (001) plane. The cations, as fourfold hydrogen-bond donors, connect neighbouring $[CI^- \cdot (H_2O)_3]_{\infty}$ anionic networks into polymeric three-dimensional hydrogenbonded assemblies.

Comment

N,N'-Bis(2-pyridyl)aryldiamines with two proton-donor and two proton-acceptor sites have been used as versatile substrates in the synthesis of extended supramolecular arrays (Bensemann *et al.*, 2002, 2003; Gdaniec *et al.*, 2002, 2005). Whereas N,N'-bis(2-pyridyl)aryldiamines cocrystallize relatively easily with dicarboxylic acids, forming good-quality crystals consisting of predictable one-dimensional hydrogenbond assemblies (Bensemann *et al.*, 2003), their salts with simple inorganic acids, with only a few exceptions, are difficult to crystallize. Here, we report the crystal structure of 2,2'-(pphenylenediimino)dipyridinium dichloride hexahydrate, (I), which was obtained as an intermediate product in the synthesis of N,N'-di-2-pyridylbenzene-1,4-diamine (PDAB).



The asymmetric unit of (I) consists of one half of the centrosymmetric PDAB dication, one chloride ion and three water molecules (Fig. 1). In the crystalline state, PDAB has been observed in a variety of conformations, ranging from a nearly planar Z,Z form in its cocrystal with phenazine (Gdaniec *et al.*, 2005) to strongly twisted Z,Z and E,E forms in its polymorphs (Bensemann *et al.*, 2002). It was obvious that, for steric reasons, double protonation of the PDAB occurring at the pyridine N atoms should force the dication to adopt a non-planar conformation. In (I), the pirydynium and benzene units are inclined at an angle of 51.85 (9)°. They are rotated with respect to the C2/N2/C7 plane; however, as in the free base, the twist about the N2–C_{ar} bond is significantly smaller

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Figure 1

The structure of (I) with the atom-numbering scheme. Displacement ellipsoids are drawn at the 50% probability level. Labelled atoms are related to unlabelled atoms by the symmetry operation (1 - x, 2 - y, -z).

for the pyridinium ring than for the aryl unit (Table 1). The Naryl-2-pyridiniumamine group adopts a twisted Z conformation which prevents cation from binding the halide ions in a chelating mode.

The geometrical parameters of hydrogen bonds listed in Table 2 are in agreement with the values reported by Steiner (1998). The dications, which act as fourfold donors in hydrogen bonding, take part in N-H···O and N-H⁺···Cl⁻ interactions. The chloride ions and water molecules are linked via $O-H\cdots O$ and $O-H\cdots Cl^-$ hydrogen bonds into an infinite two-dimensional anionic network parallel to the (001) plane (Fig. 2a and Table 2). The cations connect the neighbouring anionic $[Cl^{-}(H_2O)_3]_{\infty}$ networks, generating a polymeric three-dimensional hydrogen-bonded assembly (Fig. 2b). Each water molecule simultaneously acts as a double donor and single acceptor in hydrogen bonding, whereas the chloride anion acts as a fivefold acceptor. The proton donors form a distorted square-pyramidal arrangement around the Cl⁻ ion, with O2W(x, y, -1 + z) situated in the apical position; the basal plane of the square pyramidal environment is open for electrostatic interaction with the electron-deficient pyridinium ring. The Cl⁻ ion is located 3.4851 (11) Å from the mean plane of the pyridinium ring at the symmetry position (1 - x, 1 - y, y)-z) and 3.54 Å from its centroid.

Experimental

The title compound, (I), was prepared as an intermediate product in the synthesis of N,N'-di-2-pyridylbenzene-1,4-diamine (Bensemann et al., 2002). The single crystal used for the X-ray analysis was obtained by recrystallization from water.



Figure 2

Hydrogen-bond interactions (dashed lines) in (I), showing (a) the twodimensional network formed by Cl^- anions and water molecules and (b)the crystal packing viewed along the *a* axis, with H atoms omitted for clarity.

Crystal data $C_{16}H_{16}N_4^{2+} \cdot 2Cl^{-} \cdot 6H_2O$ $V = 555.14 (14) \text{ Å}^3$ $M_r = 443.32$ Z = 1Triclinic, P1 $D_{\rm r} = 1.326 {\rm Mg} {\rm m}^{-3}$ a = 8.2870 (10) ÅMo $K\alpha$ radiation $\mu = 0.33 \text{ mm}^{-1}$ b = 8.3104 (10) ÅT = 170 (2) K c = 9.1906 (12) Å $\alpha = 65.572 (12)^{\circ}$ Block, colourless $\beta = 89.030 (10)^{\circ}$ $0.40 \times 0.40 \times 0.40$ mm

Data collection

 $\gamma = 75.472 (10)^{\circ}$

Kuma KM4CCD κ-geometry	1942 independent reflections
diffractometer	1757 reflections with $I > 2\sigma(I)$
ω scans	$R_{\rm int} = 0.027$
Absorption correction: none	$\theta_{\rm max} = 25.0^{\circ}$
3011 measured reflections	

Refinement

Refinement on F^2 $w = 1/[\sigma^2(F_o^2) + (0.0572P)^2]$ $R[F^2 > 2\sigma(F^2)] = 0.036$ + 0.1307P] $wR(F^2) = 0.101$ where $P = (F_0^2 + 2F_c^2)/3$ S = 1.07 $(\Delta/\sigma)_{\rm max} < 0.001$ $\Delta \rho_{\rm max} = 0.21 \ {\rm e} \ {\rm \AA}$ 1942 reflections 142 parameters $\Delta \rho_{\rm min} = -0.22 \text{ e } \text{\AA}^{-3}$ Only H-atom displacement para-Extinction correction: SHELXL97 meters refined Extinction coefficient: 0.064 (10)

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 $I > 2\sigma(I)$

Table 1	
Selected geometric parameters (Å, °).	

N1-C2 N1-C6	1.343 (2) 1.362 (2)	N2-C2 N2-C7	1.338 (2) 1.4264 (19)	
C2-N1-C6	122.08 (15)	C2-N2-C7	128.24 (13)	
C7-N2-C2-N1	-13.8 (3)	C2-N2-C7-C9	-45.1 (2)	

Table 2

Hydrogen-bond geometry (Å, °).

$D - H \cdots A$	D-H	$H \cdot \cdot \cdot A$	$D \cdot \cdot \cdot A$	$D - H \cdots A$
$O1W-H1WA\cdots O2W$	0.85	1.94	2.779 (2)	168
$O1W - H1WB \cdots O3W$	0.85	1.91	2.759 (2)	175
$O2W-H2WA\cdots Cl1^{i}$	0.85	2.35	3.1957 (17)	175
O2W−H2WB···Cl1 ⁱⁱ	0.85	2.37	3.2067 (16)	168
O3W−H3WA···Cl1 ⁱⁱⁱ	0.85	2.36	3.1986 (17)	169
O3W−H3WB···Cl1 ^{iv}	0.85	2.39	3.2036 (17)	161
$N1-H1\cdots O1W$	0.90	1.85	2.7121 (18)	160
$N2-H2\cdots Cl1$	0.90	2.31	3.2056 (15)	179

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

All H atoms were located in electron-density difference maps. H atom distances were standardized to 0.85, 0.90 and 0.96 Å for O-H,

N-H and C-H bonds, respectively. H atoms were refined as riding on their parent atoms with freely refined U_{iso} values.

Data collection: *CrysAlis CCD* (Oxford Diffraction, 2000); cell refinement: *CrysAlis CCD*; data reduction: *CrysAlis RED* (Oxford Diffraction, 2000); program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *Stereochemical Workstation Operation Manual* (Siemens, 1989); software used to prepare material for publication: *SHELXL97*.

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References

Bensemann, I., Gdaniec, M., Łakomecka, K., Milewska, M. J. & Połoński, T. (2003). Org. Biomol. Chem. 1, 1425–1434.

Bensemann, I., Gdaniec, M. & Połoński, T. (2002). *New J. Chem.* **26**, 448–456. Gdaniec, M., Bensemann, I. & Połoński, T. (2002). *Acta Cryst.* **C58**, 0735–0738. Gdaniec, M., Bensemann, I. & Połoński, T. (2005). *CrystEngComm*, **7**, 433–

438. Oxford Diffraction (2000). *CrysAlis CCD* and *CrysAlis RED*. Versions 1.163. Oxford Diffraction, Abingdon, Oxfordshire, England.

Sheldrick, G. M. (1997). SHELXL97 and SHELXS97. University of Götingen, Germany.

Siemens (1989). Stereochemical Workstation Operation Manual. Release 3.4. Siemens Analytical X-ray Instruments Inc., Madison, Wisconsin, USA. Steiner, T. (1998). Acta Cryst. B54, 456–463.