

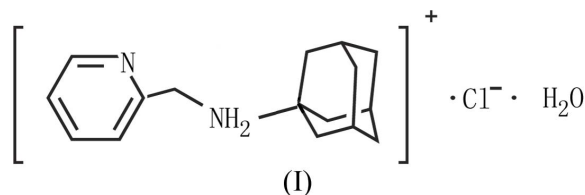
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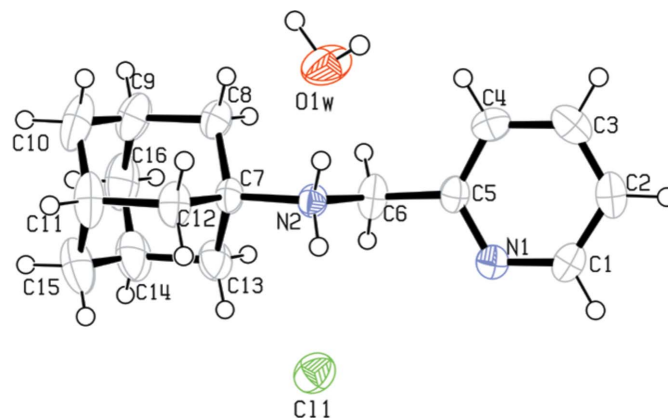
## Key indicators

Single-crystal X-ray study  
 $T = 293$  K  
Mean  $\sigma(\text{C}-\text{C}) = 0.002$  Å  
 $R$  factor = 0.041  
 $wR$  factor = 0.131  
Data-to-parameter ratio = 21.5For details of how these key indicators were  
automatically derived from the article, see  
<http://journals.iucr.org/e>.***N*-(2-Pyridylmethyl)adamantane-1-ammonium  
chloride monohydrate**In the crystal structure of the title compound,  $\text{C}_{16}\text{H}_{23}\text{N}_2^+\cdot\text{Cl}^-\cdot\text{H}_2\text{O}$ , the molecules and ions are linked through  $\text{N}-\text{H}\cdots\text{Cl}$ ,  $\text{O}-\text{H}\cdots\text{Cl}$ ,  $\text{O}-\text{H}\cdots\text{N}$  and  $\text{N}-\text{H}\cdots\text{O}$  hydrogen bonds. In addition,  $\pi-\pi$  stacking between pyridine rings leads to a tail-to-tail arrangement.Received 8 November 2006  
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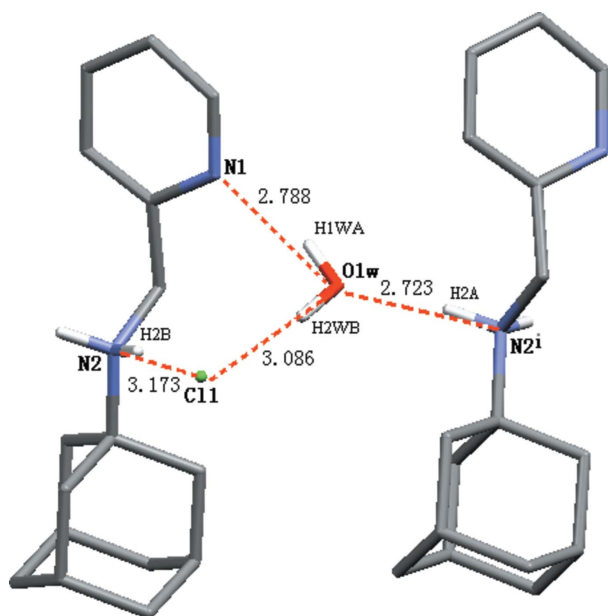
## Comment

Recently, a relatively new family of macrocyclic compounds, the cucurbit[*n*]urils ((Freeman *et al.*, 1981; Day & Arnold, 2000; Day *et al.*, 2002; Kim *et al.*, 2000) and their derivatives (Zhao *et al.*, 2001; Isobe *et al.*, 2002; Jon *et al.*, 2003; Day *et al.*, 2003) or analogues (Lagona, Fettingner & Isaacs, 2005; Wagner *et al.*, 2005), have been studied extensively in molecular recognition, host-guest chemistry and supramolecular chemistry, and summarized in a number of reviews (Lagona, Mukhopadhyay *et al.*, 2005; Lee *et al.*, 2003; Gerasko *et al.*, 2002). In the present report, we introduced a guest with two units that could each interact with various cucurbit[*n*]urils, namely *N*-(2-pyridylmethyl)adamantane-1-amine.

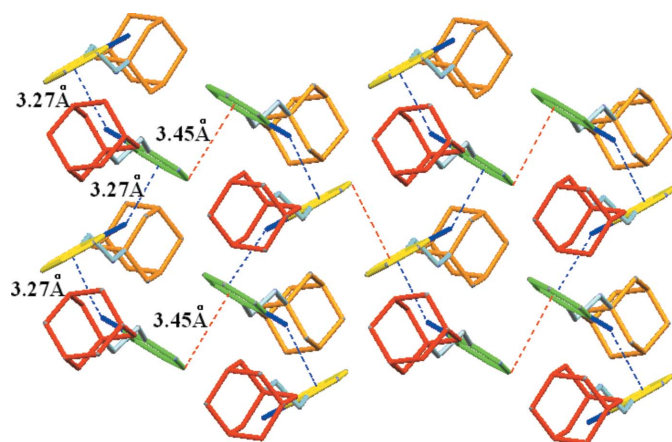
In the crystal structure of the title compound, (I), water molecule OW1 forms hydrogen bonds with the amine nitrogen on the pyridyl, N1, the protonated amine nitrogen N2 of an adjacent cation and the counter-ion C11 (see Table 1 and



**Figure 1**  
The molecular structure of (I), showing the atom-labelling scheme. Displacement ellipsoids are drawn at the 50% probability level.



**Figure 2**  
The hydrogen-bonding scheme. Hydrogen bonds are shown as dashed lines. H atoms not involved in hydrogen bonding have been omitted. [Symmetry code: (i)  $\frac{1}{2} + x, \frac{1}{2} - y, z$ .]



**Figure 3**  
Packing diagram of (I).  $\pi$ - $\pi$  Stacking and C-H... $\pi$  interactions are shown as red and blue dashed lines, respectively. H atoms have been omitted for clarity.

Fig. 2). In addition, the protonated amine nitrogen N2 forms a further hydrogen bond with C11.

Layers are formed with cations packing in a tail-to-tail manner (Fig. 3, pyridine units in yellow and green), such that the adamantaneamine units (in red and orange) are far away from each other. The driving force for this self-assembled structure can be attributed to  $\pi$ - $\pi$  stacking and C-H... $\pi$  interactions (Lin *et al.*, 2003; Tao *et al.*, 2003, 2004) between the aromatic pyridyl rings in the title compound.  $\pi$ - $\pi$  Stacking occurs between adjacent pyridine rings of the same colour (green to green or yellow to yellow); the distance (red dashed lines) between two rings is about 3.27 Å (Fig. 3). The C-H... $\pi$  interaction occurs between adjacent pyridine rings of

different colours, for example, the carbon with the blue proton on a green pyridine ring points to the top of an adjacent yellow pyridine ring; the C-H... $\pi$  distance (blue dashed lines) is about 3.45 Å. Thus, the crystal structure of the title compound is built up through a combination of hydrogen bonds,  $\pi$ - $\pi$  stacking and C-H... $\pi$  interactions.

## Experimental

A solution of 2-pyridinecarbaldehyde (1.5 ml, 8.92 mmol) in  $\text{CHCl}_3$  (10 ml) was added to a stirred solution of 1-adamantaneamine (1.41 g, 8.92 mmol) in  $\text{CHCl}_3$  (40 ml) on an ice bath over a period of 10 min.  $\text{MgSO}_4$  (1 g) was then added to the mixture. The  $\text{MgSO}_4$  was removed by filtration after 1.5 h, and the solvent was removed by evaporation, yielding a solid material, which was dissolved in ethanol (20 ml). To this stirred solution, a solution of  $\text{NaBH}_4$  (0.678 g) in ethanol (20 ml) was added dropwise at 273 K for 2 h and then filtered. The filtrate was neutralized with HCl to pH 6–7 and filtered again. Concentrated HCl (10 ml) was added to the filtrate containing (I), which was left to stand in a cool room overnight. 1.5 g of colourless crystals were collected, in a yield of 62%.

### Crystal data

$\text{C}_{16}\text{H}_{23}\text{N}_2^+\cdot\text{Cl}^-\cdot\text{H}_2\text{O}$   
 $M_r = 296.83$   
 Monoclinic,  $P2_1/a$   
 $a = 12.6019$  (6) Å  
 $b = 7.1780$  (4) Å  
 $c = 17.6949$  (9) Å  
 $\beta = 94.783$  (3)°  
 $V = 1595.04$  (14) Å<sup>3</sup>

$Z = 4$   
 $D_x = 1.236$  Mg m<sup>-3</sup>  
 Mo  $K\alpha$  radiation  
 $\mu = 0.24$  mm<sup>-1</sup>  
 $T = 293$  (2) K  
 Prism, colourless  
 $0.21 \times 0.18 \times 0.15$  mm

### Data collection

Bruker SMART CCD area-detector diffractometer  
 $\varphi$  and  $\omega$  scans  
 Absorption correction: multi-scan (SADABS; Bruker, 2002)  
 $T_{\min} = 0.952$ ,  $T_{\max} = 0.965$

14934 measured reflections  
 3962 independent reflections  
 3072 reflections with  $I > 2\sigma(I)$   
 $R_{\text{int}} = 0.026$   
 $\theta_{\max} = 28.3^\circ$

### Refinement

Refinement on  $F^2$   
 $R[F^2 > 2\sigma(F^2)] = 0.041$   
 $wR(F^2) = 0.131$   
 $S = 1.08$   
 3962 reflections  
 184 parameters  
 H-atom parameters constrained

$w = 1/[\sigma^2(F_o^2) + (0.0693P)^2 + 0.2198P]$   
 where  $P = (F_o^2 + 2F_c^2)/3$   
 $(\Delta/\sigma)_{\max} = 0.001$   
 $\Delta\rho_{\max} = 0.30$  e Å<sup>-3</sup>  
 $\Delta\rho_{\min} = -0.20$  e Å<sup>-3</sup>  
 Extinction correction: SHELXL97  
 Extinction coefficient: 0.012 (2)

**Table 1**

Hydrogen-bond geometry (Å, °).

D—H...A	D—H	H...A	D...A	D—H...A
N2—H2A...O1W	0.90	1.83	2.7228 (15)	170
N2—H2B...C11	0.90	2.28	3.1729 (13)	171
O1W—H1WA...N1 <sup>i</sup>	0.86	1.96	2.7885 (16)	162
O1W—H2WB...C11 <sup>i</sup>	0.94	2.16	3.0859 (13)	168

Symmetry code: (i)  $x + \frac{1}{2}, -y + \frac{1}{2}, z$ .

Water H atoms were located in a difference Fourier map and refined in their as-found positions relative to the O atoms with  $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{O})$ . Other H atoms were placed in calculated

positions, with C–H = 0.93–0.98 Å and N–H = 0.90 Å, and refined using a riding model, with  $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C,N})$ .

Data collection: *SMART* (Bruker, 2002); cell refinement: *SAINTE* (Bruker, 2002); data reduction: *SAINTE*; 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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