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## Key indicators

Single-crystal X-ray study
$T=293 \mathrm{~K}$
Mean $\sigma(\mathrm{C}-\mathrm{C})=0.002 \AA$
$R$ factor $=0.032$
$w R$ factor $=0.070$
Data-to-parameter ratio $=18.0$
For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e.
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## 4-Acetylanilinium chloride monohydrate

The crystal structure of the title compound, $\mathrm{C}_{8} \mathrm{H}_{10} \mathrm{NO}^{+} \cdot \mathrm{Cl}^{-}$.$\mathrm{H}_{2} \mathrm{O}$, is composed of 4-acetylanilinium cations, chloride anions and water molecules of crystallization. The crystal packing features stacking interactions between the aromatic rings, with stacks running along the $c$ axis of the crystal, as well as extensive $\mathrm{N}-\mathrm{H} \cdots \mathrm{O}, \mathrm{N}-\mathrm{H} \cdots \mathrm{Cl}$ and $\mathrm{O}-\mathrm{H} \cdots \mathrm{Cl}$ hydrogen bonding which involves all five 'active' H atoms and links cations, anions and water molecules into a three-dimensional framework.

## Comment

The title salt, (I), is composed of 4-acetylanilinium $\left(\mathrm{C}_{8} \mathrm{H}_{10} \mathrm{NO}^{+}\right)$cations, chloride anions and water molecules of crystallization. The cation is a protonated aniline derivative (Fig. 1); protonation results in a lengthening of the $\mathrm{C} 1-\mathrm{N} 1$ bond to 1.454 (2) $\AA$. This bond is significantly longer than that in non-protonated anilines [1.335 (3) Å (Goswami et al., 1999), 1.386 (4) $\AA$ (Ploug-Sørensen \& Andersen, 1985) and 1.391 (3) $\AA$ (Ploug-Sørensen \& Andersen, 1986)] and is typical for anilinium cations [1.464 (2) $\AA$ (Ploug-Sørensen \& Andersen, 1985) and 1.460 (2) $\AA$ (Ploug-Sørensen \& Andersen, 1986)].

(I)

The most significant feature of the crystal packing is the stack of aromatic rings running along the $c$ axis of the crystal; the distances between the centroids of neighbouring rings within the stack are 3.647 (2) and 3.648 (2) $\AA$ (Fig. 2). The chloride anions occupy the channels between the stacks.

All five 'active' H atoms in the structure are involved in hydrogen bonds of the $\mathrm{N}-\mathrm{H} \cdots \mathrm{O}, \mathrm{O}-\mathrm{H} \cdots \mathrm{Cl}$ and $\mathrm{N}-\mathrm{H} \cdots \mathrm{Cl}$ types (Table 2), linking all entities present in the structure into an infinite three-dimensional framework.

Figure 1


An ORTEP-3 (Farrugia, 1997) view of the cation, anion and water molecule in the structure of the title compound, showing the atomlabelling scheme and displacement ellipsoids at the $50 \%$ probability level. Hydrogen bonds are shown as dashed lines.

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Figure 2
View of the packing of the title compound. Hydrogen bonds are shown as dashed lines.

## Experimental

The title compound, (I), was prepared by bubbling gaseous hydrogen chloride through a solution of 4-acetylaniline ( 2 g ) in ether ( 50 ml ) for 15 min at an approximate flow rate of $80 \mathrm{ml} \mathrm{min}^{-1}$. The precipitate was filtered off and recrystallized from a mixture of ethanol and concentrated hydrochloric acid. The recrystallized product was dried in a desiccator, first over potassium hydroxide and then over phosphorus pentoxide.

## Crystal data

## $\mathrm{C}_{8} \mathrm{H}_{10} \mathrm{NO}^{+} \cdot \mathrm{Cl}^{-} \cdot \mathrm{H}_{2} \mathrm{O}$ <br> $M_{r}=189.64$ <br> Monoclinic, $C c$ <br> $a=7.690$ (1) $\AA$ <br> $b=17.930(1) \AA$ <br> $c=7.095$ (1) $\AA$ <br> $\beta=95.70(1)^{\circ}$ <br> $V=973.4(2) \AA^{3}$ <br> $Z=4$

$D_{x}=1.294 \mathrm{Mg} \mathrm{m}^{-3}$
Mo $K \alpha$ radiation
Cell parameters from 9241
$\quad$ reflections
$\theta=2.3-28.9^{\circ}$
$\mu=0.35 \mathrm{~mm}^{-1}$
$T=293(2) \mathrm{K}$
Prism, light yellow
$0.30 \times 0.21 \times 0.21 \mathrm{~mm}$

## Data collection

Stoe IPDS-II diffractometer $\omega$ scans
Absorption correction: numerical
( $X$-RED32; Stoe \& Cie, 2002)
$T_{\text {min }}=0.915, T_{\max }=0.928$
7188 measured reflections
2447 independent reflections

## Refinement

Refinement on $F^{2}$
$R\left[F^{2}>2 \sigma\left(F^{2}\right)\right]=0.032$
$w R\left(F^{2}\right)=0.070$
$S=0.96$
2447 reflections
136 parameters
H atoms treated by a mixture of independent and constrained refinement

Table 1
Selected geometric parameters $\left(\AA^{\circ},^{\circ}\right)$.

| $\mathrm{C} 1-\mathrm{N} 1$ | $1.454(2)$ | $\mathrm{C} 7-\mathrm{O} 1$ | $1.211(2)$ |
| :--- | :--- | :--- | ---: |
|  |  |  |  |
| $\mathrm{C} 2-\mathrm{C} 1-\mathrm{N} 1$ | $119.4(1)$ | $\mathrm{O} 1-\mathrm{C} 7-\mathrm{C} 8$ | $121.5(2)$ |
| $\mathrm{C} 6-\mathrm{C} 1-\mathrm{N} 1$ | $118.74(13)$ | $\mathrm{C} 4-\mathrm{C} 7-\mathrm{C} 8$ | $118.8(2)$ |
| $\mathrm{O} 1-\mathrm{C} 7-\mathrm{C} 4$ | $119.7(1)$ |  |  |

Table 2
Hydrogen-bonding geometry $\left(\AA{ }^{\circ},{ }^{\circ}\right)$.

| $D-\mathrm{H} \cdots A$ | $D-\mathrm{H}$ | $\mathrm{H} \cdots A$ | $D \cdots A$ | $D-\mathrm{H} \cdots A$ |
| :--- | :--- | :--- | :--- | :--- |
| $\mathrm{~N} 1-\mathrm{H} 1 A \cdots \mathrm{O}^{\mathrm{i}}$ | 0.89 | 1.94 | $2.826(2)$ | 175 |
| $\mathrm{O}^{\mathrm{i}} W-\mathrm{H} 1 W \cdots \mathrm{Cl} 1^{\mathrm{ii}}$ | $0.85(4)$ | $2.32(4)$ | $3.147(2)$ | $165(3)$ |
| $\mathrm{O} 1 W-\mathrm{H} 2 W \cdots \mathrm{Cl} 1^{\text {iii }}$ | $0.88(3)$ | $2.21(3)$ | $3.084(2)$ | $174(3)$ |
| $\mathrm{N} 1-\mathrm{H} 1 B \cdots \mathrm{Cl} 1$ | 0.89 | 2.21 | $3.089(2)$ | 171 |
| $\mathrm{~N} 1-\mathrm{H} 1 C \cdots \mathrm{O} 1 W$ | 0.89 | 1.81 | $2.697(2)$ | 175 |

Symmetry codes: (i) $x-1,1-y, z-\frac{1}{2}$; (ii) $x, y, 1+z$; (iii) $x-\frac{1}{2}, \frac{3}{2}-y, \frac{1}{2}+z$.

The aromatic and water H atoms were refined isotropically. In the aromatic ring, the refined $\mathrm{C}-\mathrm{H}$ distances are in the range 0.92 (2)0.98 (2) A. In the water molecule, the $\mathrm{O}-\mathrm{H}$ distances are 0.85 (4) and 0.88 (3) $\AA$. The methyl and anilinium N -bound H atoms were included in the refinement in a riding-model approximation, with C $\mathrm{H}=0.96 \AA, \mathrm{~N}-\mathrm{H}=0.89 \AA$ and $U_{\text {iso }}=1.5 U_{\text {eq }}$ (carrier atom).

Data collection: $X$-AREA (Stoe \& Cie, 2002); cell refinement: $X$-AREA; data reduction: $X$-RED32 (Stoe \& Cie, 2002); program(s) used to solve structure: SHELXS97 (Sheldrick, 1990); 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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