

## 2,5-Dicarboxyanilinium chloride monohydrate

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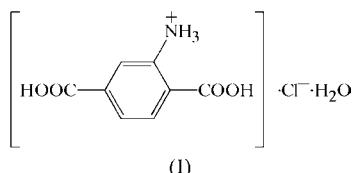
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The title compound,  $C_8H_8NO_4^+\cdot Cl^- \cdot H_2O$ , is the chlorohydrated form of 2-aminobenzene-1,4-dicarboxylic acid, the basic crystal structure of which is still not known. Molecules are linked by classical N–H $\cdots$ O, O–H $\cdots$ O, N–H $\cdots$ Cl and O–H $\cdots$ Cl hydrogen bonds, mainly along the molecular plane, into sheets built by unusual  $R_6^4(26)$ ,  $R_6^4(22)$  and  $R_4^3(22)$  rings. The stacking between layers is stabilized by another N–H $\cdots$ Cl hydrogen bond and by  $\pi$ – $\pi$  interactions between aromatic rings facing each other.

### Comment

It is known that several aromatic species with functional groups develop specific intermolecular interactions, providing tectons (specific building blocks; Hosseini, 2003) suitable for crystal engineering. In particular, aromatic carboxylic acid (Biradha *et al.*, 1998; Ferguson *et al.*, 1998; Félix *et al.*, 2000) and amino groups (Bensemann *et al.*, 2003; Karle *et al.*, 2003) can act as hydrogen-bond donors and acceptors, keeping the direction of the hydrogen bonds approximately along the molecular plane. Often, the flat extension of such hydrogen-bond interactions leads to one-dimensional supramolecular



tapes or two-dimensional supramolecular layers (Russell & Ward, 1996), whereas  $\pi$ – $\pi$  stacking interactions will link layers into a three-dimensional framework (Glidewell *et al.*, 2004). These substrates are widely harnessed in the design of functional organic solids, and also in the building of metal–organic scaffolds with specific properties (Shi *et al.*, 2005; Arora & Pedireddi, 2003). 2-Aminobenzene-1,4-dicarboxylic acid

(2-aminoterephthalic acid, 2aTPT) is well known as a ligand for metal–organic frameworks (MOFs) for systematic structural design, with possible applications in methane storage (Eddaoudi *et al.*, 2002). Because of its interesting functional groups, in order to extend the  $\pi$ -conjugated system we have

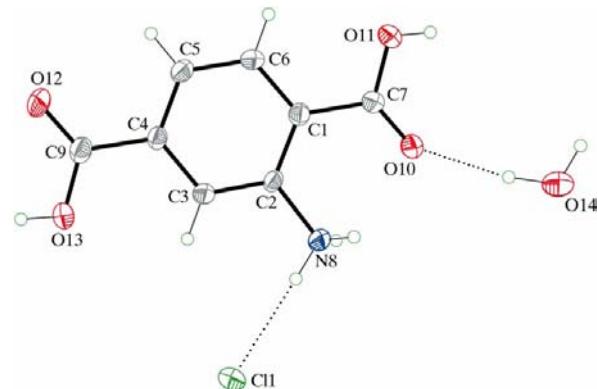


Figure 1

A view of (I), showing the atom-numbering scheme. Displacement ellipsoids are drawn at the 30% probability level and H atoms are shown as small spheres of arbitrary radii. Dotted lines indicate hydrogen bonds.

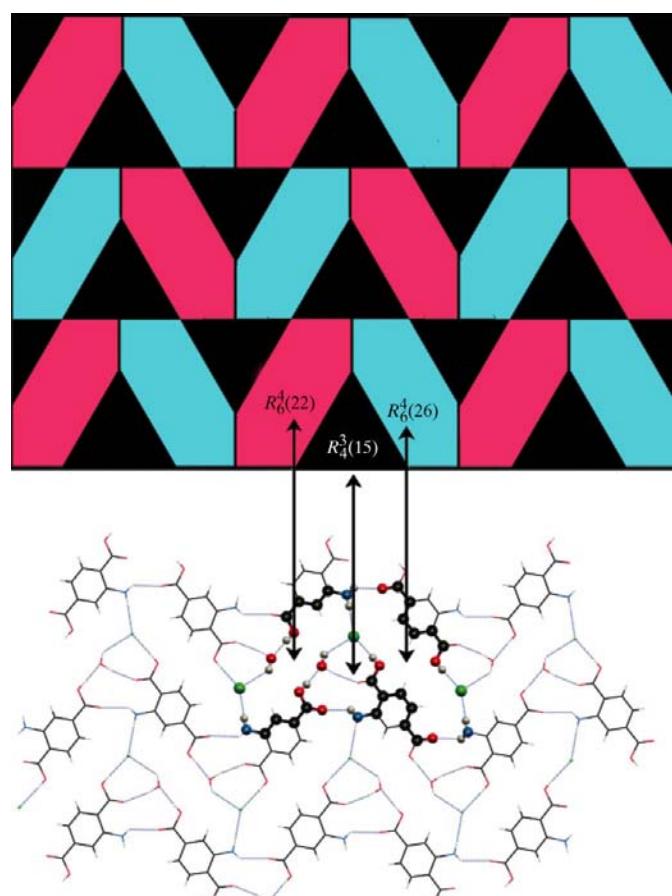


Figure 2

A perpendicularly oriented view of the (201) crystallographic plane covered by  $R_6^4(26)$ ,  $R_6^4(22)$  and  $R_4^3(15)$  rings, whose disposition is shown also in the schematic representation. Dotted lines indicate hydrogen bonds.

obtained the corresponding salicylideneimine by a condensation reaction (unpublished data). Starting from simple structural reports concerning directional intermolecular interactions (Bruno *et al.*, 2004), we decided to move on to the study of chlorohydrated 2aTPT, (I), since suitable crystals were obtained from water–methanol solutions.

Compound (I) crystallizes in the  $P2_1/c$  space group. The asymmetric unit contains one 2,5-dicarboxyanilinium cation, a  $\text{Cl}^-$  counter-ion and a water molecule. The aromatic cation presents a planar conformation [maximum deviation from the mean molecular plane for atom O2 of 0.108 (5) Å; see also geometric parameters in Table 1].

The crystal packing of (I) is mainly stabilized by classical hydrogen bonds involving the water molecule and the  $\text{Cl}^-$  anion (Table 2). As the O–H and N–H groups are hydrogen-bond donors along the molecular plane, linked to coplanar acceptors, the development of a corrugated supramolecular layer is observed. In this bidimensional array lying roughly along the (201) crystallographic plane, several repeating ring units, determined by the above-mentioned hydrogen bonds, can be distinguished. The surface is covered by two centrosymmetric hexagon-shaped units, with graph-set motifs (Bernstein *et al.*, 1995) of  $R_6^4(26)$  and  $R_6^3(22)$ , and by a triangular ring, with graph-set  $R_4^3(15)$  (Fig. 2). Along the perpendicular direction, the undulating sheets are stacked by a vertical hydrogen bond [ $\text{N}8 \cdots \text{Cl}1^i = 3.229$  (3) Å,  $\text{H}8C \cdots \text{Cl}1^i = 2.35$  Å and  $\text{N}8 - \text{H}8C \cdots \text{Cl}1^i = 167^\circ$ ; symmetry code: (i)  $x, -y + \frac{3}{2}, z - \frac{1}{2}$ ] and also by  $\pi-\pi$  interactions; the distance between close aromatic rings facing each other is 3.374 Å.

## Experimental

2aTPT (4 mg) was dissolved in hot methanol (5 ml) and then 1 M aqueous hydrochloric acid (35 µl) was added. Brown crystals of (I) suitable for X-ray analysis were obtained after slow evaporation of the resulting mixture.

### Crystal data

$\text{C}_8\text{H}_8\text{NO}_4^+ \cdot \text{Cl}^- \cdot \text{H}_2\text{O}$	$Z = 4$
$M_r = 235.62$	$D_x = 1.591 \text{ Mg m}^{-3}$
Monoclinic, $P2_1/c$	Mo $K\alpha$ radiation
$a = 6.5170$ (10) Å	$\mu = 0.39 \text{ mm}^{-1}$
$b = 16.670$ (3) Å	$T = 298$ (2) K
$c = 9.560$ (2) Å	Irregular, brown
$\beta = 108.69$ (2)°	$0.40 \times 0.19 \times 0.11$ mm
$V = 983.8$ (3) $\text{\AA}^3$	

### Data collection

Bruker P3 diffractometer	1155 reflections with $I > 2\sigma(I)$
$\omega/2\theta$ scans	$R_{\text{int}} = 0.024$
Absorption correction: $\psi$ scan (North <i>et al.</i> , 1968)	$\theta_{\text{max}} = 25.1^\circ$
$T_{\text{min}} = 0.860$ , $T_{\text{max}} = 0.962$	1 standard reflection
2395 measured reflections	every 50 reflections
1741 independent reflections	intensity decay: none

### Refinement

Refinement on $F^2$	$w = 1/[\sigma^2(F_o^2) + (0.0478P)^2 + 0.3568P]$
$R[F^2 > 2\sigma(F^2)] = 0.043$	where $P = (F_o^2 + 2F_c^2)/3$
$wR(F^2) = 0.117$	$(\Delta/\sigma)_{\text{max}} = 0.008$
$S = 1.03$	$\Delta\rho_{\text{max}} = 0.25 \text{ e } \text{\AA}^{-3}$
1743 reflections	$\Delta\rho_{\text{min}} = -0.23 \text{ e } \text{\AA}^{-3}$
140 parameters	
H-atom parameters constrained	

**Table 1**  
Selected geometric parameters ( $\text{\AA}$ , °).

C7—O10	1.214 (3)	C9—O12	1.211 (3)
C7—O11	1.318 (3)	C9—O13	1.315 (4)
C2—C1—C7—O10	7.5 (5)	C5—C4—C9—O12	-0.1 (4)
C6—C1—C7—O10	-169.0 (3)	C3—C4—C9—O12	-179.7 (3)
C2—C1—C7—O11	-174.1 (3)	C5—C4—C9—O13	179.9 (3)
C6—C1—C7—O11	9.4 (4)	C3—C4—C9—O13	0.3 (4)

**Table 2**  
Hydrogen-bond geometry ( $\text{\AA}$ , °).

$D\cdots H\cdots A$	$D\cdots H$	$H\cdots A$	$D\cdots A$	$D\cdots H\cdots A$
N8—H8A $\cdots$ Cl1	0.89	2.3	3.186 (3)	171
N8—H8B $\cdots$ Cl1 <sup>i</sup>	0.89	2.16	2.720 (3)	120
N8—H8C $\cdots$ Cl1 <sup>ii</sup>	0.89	2.35	3.229 (3)	168
O11—H11 $\cdots$ Cl1 <sup>iii</sup>	0.82	2.19	3.002 (2)	173
O13—H13 $\cdots$ O14 <sup>iv</sup>	0.82	1.78	2.600 (3)	173

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

H atoms were located in a difference Fourier map and placed in idealized positions using the riding-model technique, with C–H = 0.93 Å, N–H = 0.89 Å and O–H = 0.82 Å, and with  $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$  or  $1.5U_{\text{eq}}(\text{N,O})$ . The best refinement was obtained with the  $\psi$ -scan (North *et al.*, 1968) absorption correction performed by *XSCANS* (Siemens, 1989).

Data collection: *P3/V* (Bruker, 1989); cell refinement: *P3/V*; data reduction: *XDISK* (Bruker, 1989); program(s) used to solve structure: *SIR2004* (Burla *et al.*, 2005); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *XP* (Bruker, 1997); software used to prepare material for publication: *PARST97* (Nardelli, 1995) and *WinGX-PC* (Version 1.6.4.05; Farrugia, 1999).

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Supplementary data for this paper are available from the IUCr electronic archives (Reference: SF3013). Services for accessing these data are described at the back of the journal.

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