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#### Key indicators

Single-crystal X-ray study  
 $T = 293$  K  
 Mean  $\sigma(\text{C}-\text{C}) = 0.005$  Å  
 $R$  factor = 0.067  
 $wR$  factor = 0.174  
 Data-to-parameter ratio = 11.0

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

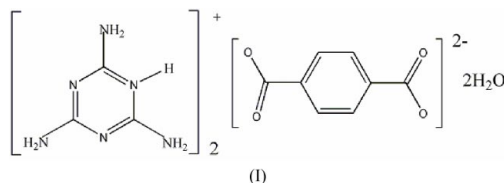
## Bis(melaminium) terephthalate dihydrate

The crystal structure of the title melaminium salt, bis(2,4,6-tri-amino-1,3,5-triazin-1-ium) terephthalate dihydrate,  $2\text{C}_3\text{H}_7\text{N}_6^{+} \cdot \text{C}_8\text{H}_4\text{O}_4^{2-} \cdot 2\text{H}_2\text{O}$ , is composed of monoprotonated melaminium cations, terephthalate dianions and water molecules. The protonated melaminium cation and the terephthalate anion are almost coplanar, the latter possessing a twofold axis of symmetry. The crystal structure involves extensive  $\text{N}-\text{H} \cdots \text{N}$ ,  $\text{N}-\text{H} \cdots \text{O}$  and  $\text{O}-\text{H} \cdots \text{O}$  hydrogen bonding.

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### Comment

We present here the crystal structure of monoprotonated bis(melaminium) terephthalate dihydrate, (I). The cation has a structure very similar to that reported for related compounds. These include:  $\text{C}_3\text{H}_7\text{N}_6^{+} \cdot \text{CH}_3\text{COO}^{-} \cdot \text{CH}_3\text{COOH} \cdot \text{H}_2\text{O}$  (Perpétuo & Janczak, 2002),  $6\text{C}_3\text{H}_7\text{N}_6^{+} \cdot \text{HPO}_4^{2-} \cdot 4\text{H}_2\text{PO}_4^{-} \cdot 4\text{H}_2\text{O}$  (Janczak & Perpétuo, 2002),  $\text{C}_3\text{H}_7\text{N}_6^{+} \cdot \text{C}_8\text{H}_5\text{O}_4^{-}$  (Janczak & Perpétuo, 2001a),  $2\text{C}_3\text{H}_7\text{N}_6^{+} \cdot \text{SO}_4^{2-} \cdot 2\text{H}_2\text{O}$  (Janczak & Perpétuo, 2001d),  $\text{C}_3\text{H}_7\text{N}_6^{+} \cdot \text{N}_3\text{O}_4^{-}$  (Tanbug *et al.*, 1999),  $\text{C}_3\text{H}_7\text{N}_6^{+} \cdot \text{NO}_3^{-}$  (Tanbug *et al.*, 1999),  $\text{C}_3\text{H}_7\text{N}_6^{+} \cdot \text{C}_4\text{H}_3\text{N}_2\text{O}_3^{-}$  (Zerkowski *et al.*, 1994),  $\text{C}_3\text{H}_7\text{N}_6^{+} \cdot \text{Cl}^{-} \cdot 0.5\text{H}_2\text{O}$  (Janczak & Perpétuo, 2001c). In addition to these monoprotonated melaminium salts, the diprotonated cation salt has also been structurally characterized in the following compounds:  $\text{C}_3\text{H}_8\text{N}_6^{2+} \cdot 2\text{C}_6\text{H}_5\text{O}_4\text{S}^{-} \cdot 2\text{H}_2\text{O}$  (Janczak & Perpétuo, 2001b) and  $\text{C}_3\text{H}_8\text{N}_6^{2+} \cdot 2\text{ClO}_4^{-} \cdot \text{H}_2\text{O}$  (Martin & Pinkerton, 1995). A search of the Cambridge Structural Database (Version 5.25; Allen, 2002) confirmed that the geometry of the terephthalate anion is normal. A twofold symmetry axis passes through the centre of the anion, bisecting the  $\text{C}5-\text{C}8^v$  and  $\text{C}8-\text{C}5^v$  bonds [symmetry code: (v)  $1-x, 2-y, 1-z$ ].



There is extensive hydrogen bonding in the structure (Fig. 2). The melaminium residues in (I) are involved in six hydrogen bonds with the anions, water and neighbouring cations. The water molecule forms two hydrogen bonds with the terephthalate (Table 1).

### Experimental

A mixture of melamine (0.126 g, 1 mmol), terephthalic acid (0.166 g, 1 mmol) and water (10 ml) was sealed in a 15 ml Teflon-lined stainless steel reactor and heated to 433 K for 60 h. Colorless crystals of the title compound suitable for X-ray analysis were obtained.

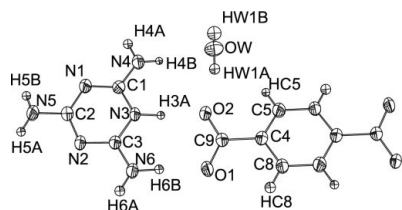


Figure 1

The molecular structure of the title compound, showing 50% probability displacement ellipsoids and the atom-numbering scheme of the asymmetric unit.

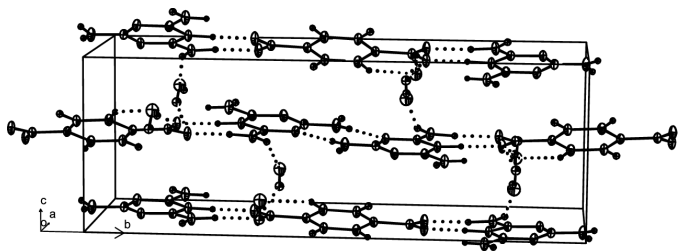


Figure 2

The packing arrangement in the unit cell, showing the hydrogen-bonding interactions as dashed lines.

#### Crystal data



$M_r = 454.44$

Monoclinic,  $P2_1/c$

$a = 6.809$  (4) Å

$b = 20.164$  (10) Å

$c = 7.062$  (6) Å

$\beta = 93.33$  (5)°

$V = 967.9$  (11) Å<sup>3</sup>

$Z = 2$

$D_x = 1.559$  Mg m<sup>-3</sup>

Mo  $K\alpha$  radiation

Cell parameters from 1132 reflections

$\theta = 2.0$ – $25.1$ °

$\mu = 0.13$  mm<sup>-1</sup>

$T = 293$  (2) K

Prism, colorless

$0.20 \times 0.16 \times 0.15$  mm

#### Data collection

Siemens SMART CCD area-detector diffractometer

$\varphi$  and  $\omega$  scans

Absorption correction: multi-scan (SADABS; Sheldrick, 1996)

$T_{\min} = 0.975$ ,  $T_{\max} = 0.981$

3095 measured reflections

1691 independent reflections

1215 reflections with  $I > 2\sigma(I)$

$R_{\text{int}} = 0.042$

$\theta_{\text{max}} = 25.1$ °

$h = -8 \rightarrow 7$

$k = -22 \rightarrow 23$

$l = -3 \rightarrow 8$

#### Refinement

Refinement on  $F^2$

$R[F^2 > 2\sigma(F^2)] = 0.067$

$wR(F^2) = 0.174$

$S = 1.08$

1691 reflections

154 parameters

H atoms treated by a mixture of independent and constrained refinement

$w = 1/[\sigma^2(F_o^2) + (0.0718P)^2 + 1.0433P]$

where  $P = (F_o^2 + 2F_c^2)/3$

$(\Delta/\sigma)_{\text{max}} = 0.170$

$\Delta\rho_{\text{max}} = 0.27$  e Å<sup>-3</sup>

$\Delta\rho_{\text{min}} = -0.26$  e Å<sup>-3</sup>

Extinction correction: SHELXL97

Extinction coefficient: 0.009 (2)

Table 1

Hydrogen-bonding geometry (Å, °).

$D-H \cdots A$	$D-H$	$H \cdots A$	$D \cdots A$	$D-H \cdots A$
$N3-H3A \cdots O2$	0.86	1.86	2.717 (4)	179
$N6-H6A \cdots OW^i$	0.86	2.18	2.959 (4)	151
$N6-H6B \cdots O1$	0.86	1.95	2.811 (4)	180
$N5-H5A \cdots N2^{ii}$	0.86	2.20	3.055 (4)	175
$N4-H4A \cdots O1^{iii}$	0.86	2.07	2.889 (4)	159
$N4-H4B \cdots OW$	0.86	2.22	3.056 (5)	163
$OW-HW1B \cdots O2$	0.87 (5)	1.88 (5)	2.738 (4)	168 (4)
$OW-HW1A \cdots O1^{iv}$	0.84 (6)	2.13 (6)	2.889 (5)	151 (6)

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

H atoms were located in a difference Fourier synthesis. The H atoms of the water molecule were refined freely. Other H atoms were refined in the riding-model approximation, with  $C-H = 0.93$  Å and  $N-H = 0.86$  Å;  $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}$  of the carrier atom.

Data collection: SMART (Siemens, 1996); cell refinement: SMART and SAINT (Siemens, 1994); data reduction: SAINT and XPREP in SHELXTL (Siemens, 1994); program(s) used to solve structure: SHELXTL; program(s) used to refine structure: SHELXTL; molecular graphics: SHELXTL; software used to prepare material for publication: SHELXTL.

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