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Melaminium bis(4-hydroxybenzenesulfonate) dihydrate

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The crystals of a new melaminium salt, 2,4,6-triamino-1,3,5triazine-1,3-diium bis(4-hydroxybenzenesulfonate) dihydrate, $C_3H_8N_6^{2+}\cdot 2C_6H_5O_4S^-\cdot 2H_2O$, are built up from doubly protonated melaminium(2+) residues, dissociated *p*-phenolsulfonate anions and water molecules. The doubly protonated melaminium dication lies on a twofold axis. The hydroxyl group of the *p*-hydroxybenzenesulfonate residue is roughly coplanar with the phenyl ring [dihedral angle 13 (2)°]. A combination of ionic and donor–acceptor hydrogen-bond interactions link the melaminium and *p*-hydroxybenzenesulfonate residues and the water molecules to form a three-dimensional network.

Comment

This study is a continuation of our investigation into the characterization of the hydrogen bonds formed by the melamine molecule in the solid state (Janczak & Perpétuo, 2001). To expand the understanding of the solid-state physicalorganic chemistry of compounds containing $N-H\cdots O$ and $O-H\cdots O$ hydrogen bonds, we have studied the solid-state structure of the title diprotonated compound, melaminium *p*-hydroxybenzenesulfonate dihydrate, (I).



and one water molecule (Fig. 1). To our knowledge, this is the third structurally characterized melaminium salt which is protonated at two ring N atoms; the first was the hydrochloride of a melaminium-cyanuric acid complex (Wang *et al.*, 1990) and the second was a hydrated complex of perchlorate acid with melamine (Martin & Pinkerton, 1995). In addition to these doubly protonated melaminium salts, singly protonated melaminium salts have also been structurally characterized (Zerkowski *et al.*, 1994; Janczak & Perpétuo, 2001).

The six-membered aromatic ring of the melaminium residue, $C_3H_8N_6^{2+}$, is almost planar. Two of the three amino groups are approximately coplanar with the weighted least-squares plane through the melaminium ring, while the third group (at N4) is rotated along the C8–N4 bond by 13 (2)°. This is probably due to an H1N1···H1N4 interaction of 2.30 (2) Å, caused by the fact that C8 is bonded to both protonated N atoms of the melaminium ring and is involved in the intermolecular hydrogen-bonding system.

The ring of the melaminium residue is significantly distorted from the ideal hexagonal form. The internal C-N-C angle at the non-protonated N atom is significantly smaller than the C-N-C angles at the protonated N atoms. These differences between the internal C-N-C angles within the melaminium ring residue correlate with the steric effect of the lone-pair electrons and are fully consistent with the valence-shell electron-pair repulsion theory (VSEPR; Gillespie, 1972). Protonation of the melamine ring at two N atoms distorts the bond lengths in the aromatic ring. The two shortest bonds in the melaminium ring (N2-C7 and its symmetrical equivalent) are those farthest from the protonated ring N atoms. The two longest N-C bonds of the melaminium ring (N1-C7 and its symmetrical equivalent) are those connected to the shortest bonds. This has the effect of opening up the ring bond angles at C7 and its symmetrical equivalent. A semi-empirical calculation performed with the AM1 parameter set (Dewar et al., 1985) on the melaminium residue doubly protonated at two ring N atoms results in almost the same geometrical features. Thus, the ring distortion of the melaminium residue mainly results from the protonation and, to a lesser degree, from the hydrogen bonding and crystal packing. The distor-





Figure 1

The molecular structure of the asymmetric unit of (I) showing 50% probability displacement ellipsoids and the atom-numbering scheme. H atoms are drawn as spheres of arbitrary radii.

tion of the aromatic melaminium ring in (I) is quite similar to that reported for the hydrochloride of the melamine-cyanuric acid complex (Wang *et al.*, 1990), as well as for the melaminium diperchlorate monohydrate complex (Martin & Pinkerton, 1995), *i.e.* both of the simple salts of diprotonated melamine that have been reported previously.

The ring of the *p*-hydroxybenzenesulfonate anion shows a slight quinone character. This is probably due to the substitution effect of the hydroxyl and sulfonate groups in the 1,4positions of the ring. The C–O (OH group) and C–S ($SO_3^$ group) bond lengths are comparable with the distances of 1.364 (15) and 1.750 (8) Å observed for $C_{aromatic}$ -O and Caromatic-S bonds, respectively (Allen et al., 1987). The hydroxyl group is roughly coplanar with the ring [C5-C4-O4-H1O4 $-13(2)^{\circ}$]. The sulfonate group has a slightly distorted tetrahedral geometry and is oriented so that the S1-O1 bond is almost coplanar with the phenyl ring [C2-C1- $S1-O1 \ 175.6 \ (1)^{\circ}$]. The differences between the S-O bond lengths of the SO₃⁻ group are correlated with the number and strength of the hydrogen bonds formed by the O atoms. The O atom of the longest S–O bond is involved in three hydrogen bonds as acceptor, while the other two O atoms are involved as acceptors in only one hydrogen bond. Although atoms O1 and O2 are involved in only one hydrogen bond, the S1-O1 and S1–O2 bond lengths are different, S1–O1 being shorter than S1–O2, since O1 forms a weaker hydrogen bond than O2.

Both oppositely charged residues and the water molecules interact extensively by a combination of ionic and donoracceptor hydrogen-bond interactions throughout the lattice to form a three-dimensional network (Fig. 2). All eight H atoms of the melaminium residue form hydrogen bonds with four different *p*-hydroxybenzenesulfonate anions and with two water molecules, which are acceptors of hydrogen bonds. Two of these four *p*-hydroxybenzenesulfonate residues are involved as acceptors in two hydrogen bonds with a melaminium residue (N1-H1N1···O2 and N4-H1N4···O3), while the other two *p*-hydroxybenzenesulfonate moieties are involved in only one hydrogen bond with the same melaminium residue. Thus, one melaminium residue forms eight hydrogen bonds. There are six N-H···O hydrogen bonds with four neighbouring *p*-hydroxybenzenesulfonate anions, and the other two H atoms of the melaminium residue form hydrogen bonds with the water molecules. The most noticeable feature is the fact that the non-protonated N atom of the melaminium residue is not involved as an acceptor in any hydrogen bond.

The SO₃⁻ group of the *p*-hydroxybenzenesulfonate residue is involved as an acceptor in three $O \cdots H - N$ hydrogen bonds from two different melaminium moieties and in two $O \cdots H - O$ hydrogen bonds with two water molecules, while the hydroxyl group of the *p*-hydroxybenzenesulfonate ion (as a donor) forms a hydrogen bond with a water molecule. Thus, one *p*-hydroxybenzenesulfonate residue is involved in six different hydrogen bonds.

The water molecule is involved as a donor in two hydrogen bonds with the SO_3^- groups of two different *p*-hydroxybenzenesulfonate anions, and as an acceptor in hydrogen bonds with the phenol hydroxyl group and with the N3 amino group from a melaminium dication.

In the crystal of (I), the melaminium residues form layers which are a/2 apart. In one layer, the melaminium residues are parallel to each other. The ring of the melaminium residue is perpendicular to the *ac* plane and forms angles of about 36 and 54° with the *bc* and *ab* planes, respectively. The ring of the *p*-hydroxybenzenesulfonate anion is almost perpendicular to the *bc* plane and makes a dihedral angle of 60.6° with the *ab* plane and 28.5° with the *ac* plane. The plane of the melaminium residue is inclined at an angle of 72.2 (1)° to the plane of the *p*-hydroxybenzenesulfonate ring. Details of the hydrogen-bonding geometry are given in Table 2.



Figure 2

The molecular arrangement in the unit cell of (I) showing the hydrogenbonding interactions (dashed lines). H atoms have been omitted for clarity.

Experimental

Melamine was dissolved in hot water and to this solution was slowly added a 10% solution of *p*-hydroxybenzenesulfonic acid. After several days, colourless crystals of (I) appeared.

Crystal data

$C_{2}H_{2}N_{2}^{2+}\cdot 2C_{2}H_{2}O_{3}S^{-}\cdot 2H_{2}O_{3}$	D measured by flotation
$M_r = 510.51$	Mo $K\alpha$ radiation
Orthorhombic, Pbcn	Cell parameters from 2155
a = 26.625 (5) Å	reflections
b = 7.863 (2) Å	$\theta = 5-26^{\circ}$
c = 10.230 (2) Å	$\mu = 0.32 \text{ mm}^{-1}$
V = 2141.7 (8) Å ³	T = 293 (2) K
Z = 4	Parallelepiped, colourless
$D_x = 1.583 \text{ Mg m}^{-3}$	$0.28 \times 0.24 \times 0.16 \text{ mm}$
$D_{\rm m} = 1.58 {\rm Mg}{\rm m}^{-3}$	

Kuma KM-4 diffractometer with a two-dimensional CCD area- detector ω scans Absorption correction: analytical, face-indexed (<i>SHELXTL</i> ; Shel- drick, 1990) $T_{\rm min} = 0.917, T_{\rm max} = 0.951$	17 563 measured reflections 2806 independent reflections 1728 reflections with $I > 2\sigma(I)$ $R_{int} = 0.034$ $\theta_{max} = 29.6^{\circ}$ $h = -36 \rightarrow 36$ $k = -10 \rightarrow 8$ $l = -14 \rightarrow 13$	
Refinement		
Refinement on F^2	$w = 1/[\sigma^2(F_o^2) + (0.0518P)^2]$	
$R[F^2 > 2\sigma(F^2)] = 0.036$	+ 0.092P]	
$wR(F^2) = 0.098$	where $P = (F_o^2 + 2F_c^2)/3$	
S = 1.06	$(\Delta/\sigma)_{\rm max} = 0.001$	
2806 reflections	$\Delta \rho_{\rm max} = 0.29 \ {\rm e} \ {\rm \AA}^{-3}$	
173 parameters	$\Delta \rho_{\rm min} = -0.27 \ {\rm e} \ {\rm \AA}^{-3}$	
H atoms treated by a mixture of	Extinction correction: SHELXL97	
independent and constrained	(Sheldrick, 1997)	

Table 1

refinement

Selected geometric parameters (Å, °).

S1-O1	1.4439 (13)	N1-C8	1.3421 (17)
S1-O2	1.4566 (13)	N1-C7	1.374 (2)
S1-O3	1.4670 (12)	N2-C7	1.3297 (18)
S1-C1	1.7528 (16)	N3-C7	1.307 (2)
O4-C4	1.355 (2)	N4-C8	1.308 (3)
01 01 02	110 50 (0)	CO 14 C7	100 56 (15)
01 - S1 - 02	112.78 (8)	C8-N1-C7	120.76 (15)
O1-S1-O3	112.37 (8)	$C7 - N2 - C7^{i}$	115.91 (18)
O2-S1-O3	110.00(7)	N3-C7-N2	120.15 (15)
O1-S1-C1	106.74 (7)	N3-C7-N1	117.08 (15)
O2-S1-C1	107.55 (7)	N2-C7-N1	122.75 (14)
O3-S1-C1	107.07 (7)	N4-C8-N1	121.5 (1)
C5-C6-C1	120.14 (14)	N1 ⁱ -C8-N1	117.0 (2)

Extinction coefficient: 0.0030 (7)

Symmetry code: (i) 1 - x, y, $\frac{3}{2} - z$.

The positions of the H atoms of the melaminium residue and of the hydroxyl (OH) group of the *p*-hydroxybenzenesulfonate ion, as well as those of the water molecule, *i.e.* of all H atoms involved in hydrogen bonding, were refined. Other H atoms were treated as riding, with C-H = 0.93 Å. For all H atoms, $U_{iso}(H) = 1.2U_{eq}(C)$ and $1.5U_{eq}(O)$.

Table 2	
Hydrogen-bonding geometry (Å	∖ , °).

$D-\mathrm{H}\cdots A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdot \cdot \cdot A$
$N1 - H1N1 \cdots O2^{i}$	0.85 (2)	1.85 (2)	2.699 (2)	172 (2)
O4-H1O4O5	0.89 (3)	1.93 (3)	2.786 (2)	162 (2)
N3-H1N3···O3 ⁱⁱ	0.86(2)	2.12 (2)	2.945 (2)	162 (2)
N3-H2N3···O5 ⁱⁱⁱ	0.86(2)	2.21(2)	2.905 (2)	138 (2)
$N4-H1N4\cdots O3^{i}$	0.93(2)	1.93 (2)	2.846 (2)	170(2)
O5−H1O5···O3 ⁱⁱⁱ	0.93(2)	2.10(2)	2.997 (2)	163 (2)
O5−H2O5···O1 ^{iv}	0.93 (2)	2.14 (2)	2.905 (2)	138 (2)

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

Data collection: *KM*-4 *CCD Software* (Kuma Diffraction, 1999); cell refinement: *KM*-4 *CCD Software*; data reduction: *KM*-4 *CCD Software*; program(s) used to solve structure: *SHELXS*97 (Sheldrick, 1997); program(s) used to refine structure: *SHELXL*97 (Sheldrick, 1997); molecular graphics: *SHELXTL* (Sheldrick, 1990); software used to prepare material for publication: *SHELXL*97.

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Supplementary data for this paper are available from the IUCr electronic archives (Reference: NA1522). Services for accessing these data are described at the back of the journal.

References

- Allen, F. M., Kennard, O., Watson, D. G., Brammer, L., Orpen, A. G. & Taylor, R. (1987). J. Chem. Soc. Perkin Trans. 2, pp. S1–19.
- Dewar, M. J. S., Zoebisch, E. G., Healy, E. F. & Stewart, J. J. P. (1985). J. Am. Chem. Soc. 107, 3902–3909.
- Gillespie, R. J. (1972). In *Molecular Geometry*. London: van Nostrand Reinhold.
- Janczak, J. & Perpétuo, G. J. (2001). Acta Cryst. C57, 123-125.
- Kuma Diffraction (1999). KM-4 CCD Software. Version 163. Kuma Diffraction. Wrocław. Poland.
- Martin, A. & Pinkerton, A. A. (1995). Acta Cryst. C51, 2174–2177.
- Sheldrick, G. M. (1990). SHELXTL. University of Göttingen, Germany.
- Sheldrick, G. M. (1997). SHELXS97 and SHELXL97. University of Göttingen, Germany.

Wang, Y., Wei, B. & Wang, Q. (1990). J. Crystallogr. Spectrosc. Res. 20, 79–84.
Zerkowski, J. A., McDonald, J. C. & Whitesides, G. M. (1994). Chem. Mater. 6, 1250–1256.