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Jian-Wei Tong, Li-Fang Ma, Deng-Ke Cao, Yi-Zhi Li and Li-Min Zheng*

Coordination Chemistry Institute, State Key Laboratory of Coordination Chemistry, Nanjing University, Nanjing 210093, People's Republic of China

Correspondence e-mail: lmzheng@nju.edu.cn

Key indicators

Single-crystal X-ray study T = 298 KMean $\sigma(\text{C}-\text{C}) = 0.005 \text{ Å}$ R factor = 0.061 wR factor = 0.161 Data-to-parameter ratio = 13.4

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

[(1*H*-Benzimidazol-2-ylmethyl)iminodimethylene]diphosphonic acid dihydrate

The title compound, $C_9H_{15}O_6N_3P_2\cdot 2H_2O$, contains two phosphonic acid groups and one benzimidazole group connected by an N(CH₂-)₃ group. One of the benzimidazole N atoms and two phosphonic acid O atoms of each PO₃ group are protonated. Extensive hydrogen-bonding interactions, as well as π - π stacking interactions, are found between the molecules, leading to a three-dimensional supramolecular network structure.

Comment

Metal phosphonates are organic-inorganic hybrid materials that are important because of their potential applications in ion exchange, sensors, catalysis and molecular recognition (Clearfield, 1998). A number of multifunctional phosphonic acids have been used for building novel structures, ranging from zero-dimensional molecules to three-dimensional frameworks. As a potential multidentate ligand, [(1H-benzimidazol-2-ylmethyl)iminodimethylene]diphosphonic acid (bibmpH₄) is potentially able to form various structures with metal ions. Five isomorphous compounds with chain structures have been obtained through hydrothermal reactions of bibmpH₄ and metal sources (Cao et al., 2006). During our efforts to synthesize a manganese compound by direct reaction of $bibmpH_4$ with $Mn(ClO_4)_2$ in aqueous solution, however, excellent quality single crystals of the unreacted acid were obtained as the title dihydrate, (I).



Fig. 1 shows the structure of compound (I) with the atomic labelling scheme. Of the three phosphonate O atoms connected to each of P1 and P2, atoms O2, O3, O5 and O6 are protonated. Hence, the P1–O2, P1–O3, P2–O5 and P2–O6 distances of 1.564 (3), 1.546 (3), 1.537 (3) and 1.567 (3) Å, respectively, are much longer than the other P1–O1 [1.509 (3) Å] and P2–O4 [1.507 (3) Å] distances. The P–C and C–O distances are normal. The CPO₃ tetrahedra are slightly distorted, with the O–P–O(C) angles ranging from 103.53 (15) to 116.84 (17)°.

Neighbouring molecules are connected by very strong hydrogen bonds (Table 1) between phosphonate atoms O2

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8348 measured reflections

 $R_{\rm int} = 0.052$

 $\theta_{\rm max} = 26.0^{\circ}$

3025 independent reflections

2151 reflections with $I > 2\sigma(I)$



Figure 1

A view of the molecular structure of (I). Displacement ellipsoids are drawn at the 30% probability level. C-bound H atoms have been omitted for clarity.



Figure 2

The crystal packing, viewed along the a axis. C-bound H atoms have been omitted for clarity. Dashed lines indicate hydrogen bonds.

and O6, forming an infinite chain along the b axis. These chains are held together through hydrogen bonds between atoms N2 and O1, and between atom O2W and the phosphonate O atoms, forming a layer in the *ab* plane.

Neighbouring layers are linked by hydrogen bonds between atoms O1W and O4 from different layers. π - π stacking interactions are also found between these layers along the a axis, with centroid-to-centroid distances between adjacent benzene rings of 3.627 (2) and 3.955 (2) Å. Therefore, a threedimensional supramolecular network is formed with onedimensional channels generated along the a axis. The solvent water molecules O1W and O2W reside within the channels.

Experimental

(Benzimidazol-2-ylmethyl)iminobis(methylenephosphonic acid) (bibmpH₄) was synthesized according to the literature procedure of Yoshikawa (1995). A mixture of bibmpH₄ (0.081 g, 0.25 mmol), manganese(II) perchlorate (0.0905 g, 0.25 mmol) and water (10 ml) was stirred at room temperature for 10 min. The filtrate was kept at room temperature for 2 d, after which colourless block crystals of (I) were collected. Analysis, calculated for (I): C 32.70, H 4.09, N 11.44%; found: C32.21, H 4.94, N 11.23%.

Crvstal data

 C_1 М Μ a b c = β v

$_{0}H_{15}N_{3}O_{6}P_{2}\cdot 2H_{2}O$	Z = 4
r = 371.22	$D_x = 1.599 \text{ Mg m}^{-3}$
onoclinic, $P2_1/c$	Mo $K\alpha$ radiation
= 7.0336 (8) Å	$\mu = 0.33 \text{ mm}^{-1}$
= 11.5822 (13) Å	T = 298 (2) K
= 18.935 (2) Å	Block, colourless
$= 91.252 (2)^{\circ}$	$0.32 \times 0.28 \times 0.26 \text{ mm}$
$= 1542.2 (3) Å^{3}$	

Data collection

- Bruker SMART APEX CCD areadetector diffractometer φ and φ scans Absorption correction: multi-scan
- (SADABS; Bruker, 2000) $T_{\min} = 0.91, T_{\max} = 0.92$

Refinement

Refinement on F^2	H-atom parameters constrained
$R[F^2 > 2\sigma(F^2)] = 0.061$	$w = 1/[\sigma^2(F_o^2) + (0.1074P)^2]$
$wR(F^2) = 0.161$	where $P = (F_0^2 + 2F_c^2)/3$
S = 0.95	$(\Delta/\sigma)_{\rm max} < 0.001$
3025 reflections	$\Delta \rho_{\rm max} = 0.66 \ {\rm e} \ {\rm \AA}^{-3}$
226 parameters	$\Delta \rho_{\rm min} = -0.64 \ {\rm e} \ {\rm \AA}^{-3}$

Table 1

Hydrogen-bond	geometry ([A, °)	
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$D - H \cdots A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdot \cdot \cdot A$
$O2W - H2WB \cdots O1W^{i}$	0.80 (7)	2.28 (7)	3.051 (5)	163 (6)
$O1W-H1WA\cdots O4^{n}$	0.85 (6)	1.98 (6)	2.823 (4)	173 (5)
O2W−H2WA····O4 ⁱⁱⁱ	0.87 (6)	1.90(7)	2.722 (5)	158 (6)
$O1W-H1WB\cdots O4^{iii}$	0.80 (6)	2.20 (6)	2.965 (4)	159 (5)
$O2-H2A\cdots O6^{iii}$	0.88 (5)	1.56 (5)	2.359 (4)	148 (4)
$O5-H5A\cdots O2W^{iv}$	0.82 (5)	2.51 (5)	3.302 (5)	162 (5)
$N2-H2C\cdotsO1^{iv}$	0.86	1.93	2.770 (4)	165
Symmetry codes: (i)	-x + 1, -y +	+1, -7 + 1;	(ii) $x_{1} - y_{2} + \frac{1}{2}$	$\frac{3}{2}$, $z = \frac{1}{2}$; (iii)

 $-x + 2, y - \frac{1}{2}, -z + \frac{3}{2};$ (iv) $-x + 1, y + \frac{1}{2}, -z + \frac{3}{2}.$

All H atoms on C and N atoms were placed in calculated positions and included as part of a riding model, with C-H = 0.93-0.97 Å and N-H = 0.86 Å, and with $U_{iso}(H) = 1.2U_{eq}$ of the parent atom. Obound H atoms were refined freely.

Data collection: SMART (Bruker, 2000); cell refinement: SAINT (Bruker, 2000); data reduction: SAINT; program(s) used to solve structure: SHELXTL (Bruker, 2000); program(s) used to refine structure: SHELXTL; molecular graphics: SHELXTL; software used to prepare material for publication: SHELXTL.

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