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Pyridinium diaguabis(methylenediphosphonato- $\kappa^2 O, O'$)chromate(III) tetrahydrate

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Key indicators: single-crystal X-ray study; T = 100 K; mean σ (C–C) = 0.013 Å; disorder in solvent or counterion; R factor = 0.024; wR factor = 0.070; data-toparameter ratio = 14.7.

In the title complex, $(C_5H_6N)[Cr(CH_4O_6P_2)_2(H_2O)_2]\cdot 4H_2O$, the Cr^{III} atom, lying on an inversion centre, is coordinated by two bidentate methylene diphosphonate ligands and two water molecules in a distorted octahedral coordination geometry. The pyridinium cation is located on an inversion centre, with an N atom and a C atom sharing a position each at a half occupancy. A three-dimensional network is constructed by $O-H\cdots O$, $N-H\cdots O$ and $C-H\cdots O$ hydrogen bonds between the pyridinium cation, complex anion and uncoordinated water molecules.

Related literature

For general background to metal-organic frameworks with diphosphonic acids, see: Barthelet et al. (2002). For related structures, see: Byun et al. (2006); Suh et al. (1997); Van der Merwe et al. (2009); Visser et al. (2010).



Experimental

Crystal data	
$(C_5H_6N)[Cr(CH_4O_6P_2)_2(H_2O)_2]$	$\beta = 106.128 \ (5)^{\circ}$
$4H_2O$	$\gamma = 94.496 \ (5)^{\circ}$
$M_r = 588.17$	V = 535.7 (6) Å ³
Triclinic, P1	Z = 1
a = 7.206 (5) Å	Mo $K\alpha$ radiation
b = 7.485(5) Å	$\mu = 0.92 \text{ mm}^{-1}$
c = 10.984 (5) Å	T = 100 K
$\alpha = 107.085(5)^{\circ}$	$0.22 \times 0.16 \times 0.08 \text{ mm}$

 $R_{\rm int} = 0.020$

8784 measured reflections

2632 independent reflections

2483 reflections with $I > 2\sigma(I)$

Data collection

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Bruker APEXII CCD
  diffractometer
Absorption correction: multi-scan
  (SADABS; (Bruker, 2001)
  T_{\min} = 0.843, T_{\max} = 0.931
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Refinement

$R[F^2 > 2\sigma(F^2)] = 0.024$	H atoms treated by a mixture of
$wR(F^2) = 0.070$	independent and constrained
S = 1.05	refinement
2632 reflections	$\Delta \rho_{\rm max} = 0.47 \ {\rm e} \ {\rm \AA}^{-3}$
179 parameters	$\Delta \rho_{\rm min} = -0.62 \text{ e } \text{\AA}^{-3}$
16 restraints	

Table 1

Selected bond lengths (Å).

Cr1-O1 Cr1-O2	1.991 (4) 1.956 (4)	Cr1-07	1.964 (4)
011 02	1000(1)		

Table 2

Hydrogen-bon	d geometry	(Å, °`).

$D - H \cdot \cdot \cdot A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdots A$
$C1 - H4 \cdots O6^{i}$	0.97	2.49	3.346 (7)	147
$C4-H4A\cdots O9^{ii}$	0.93	2.16	2.93 (7)	140
$N1 - H1 \cdots O9^{ii}$	0.86	2.32	3.03 (5)	141
$O1-H1A\cdots O6^{i}$	0.80 (6)	1.83 (6)	2.634 (6)	176 (9)
$O1-H1B\cdots O4^{iii}$	0.83 (6)	1.87 (6)	2.704 (6)	177 (9)
$O3-H3\cdots O8^{iv}$	0.82	1.83	2.629 (6)	163
O5−H6···O4 ⁱⁱ	0.83 (5)	1.80 (5)	2.619 (6)	175 (10)
$O8-H7\cdots O6^{v}$	0.83 (6)	1.86 (6)	2.687 (6)	171 (9)
O8−H8···O9	0.85 (7)	1.94 (8)	2.748 (7)	158 (11)
$O9-H9A\cdots O4$	0.83 (6)	2.00 (6)	2.833 (6)	179 (10)
$O9-H10\cdots O8^{vi}$	0.84 (7)	1.99 (7)	2.820 (7)	174 (13)

Symmetry codes: (i) -x + 1, -y + 2, -z + 1; (ii) -x, -y + 1, -z + 1; (iii) x + 1, y, z; (iv) -x + 1, -y + 2, -z + 2; (v) x, y, z + 1; (vi) -x, -y + 2, -z + 2.

Data collection: APEX2 (Bruker, 2007); cell refinement: SAINT-Plus (Bruker, 2007); data reduction: SAINT-Plus; program(s) used to solve structure: SHELXTL (Sheldrick, 2008); program(s) used to refine structure: SHELXTL; molecular graphics: DIAMOND (Brandenburg, 1999); software used to prepare material for publication: SHELXTL.

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: HY2333).

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supporting information

Acta Cryst. (2010). E66, m1011–m1012 [https://doi.org/10.1107/S1600536810028990] Pyridinium diaquabis(methylenediphosphonato-κ²O,O')chromate(III) tetrahydrate

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S1. Comment

The title compound forms part of an ongoing study in our group involving methylene diphosphonate and its coordination to various metal cores. (Van der Merwe *et al.*, 2009; Visser *et al.*, 2010). Diphosphonic acids are useful for the synthesis of metal-organic frameworks exhibiting microporous properties (Barthelet *et al.*, 2002).

The Cr^{III} ion in the title complex is in a distorted octahedral environment (Fig. 1), with Cr—O bond distances ranging from 1.956 (4) to 1.991 (4) Å (Table 1). All the bond distances and angles are well within the normal range (Byun *et al.*, 2006; Suh *et al.*, 1997). The pyridinium cation is located on an inversion centre and an N atom and a C atom share a position at a half occupancy for each atom. A three-dimensional network is provided by numerous hydrogen bonds between the pyridinium cation, complex anion and uncoordinated water molecules (Table 2).

S2. Experimental

 $CrCl_{3.6}H_{2}O$ (0.092 g, 0.347 mmol) was dissolved in water (40 ml) and ammonium hydroxide was gradually added dropwise in order to precipitate Cr(III) hydroxide. Methylene diphosphonate (0.347 g, 2 mmol) was added to the $Cr(OH)_{3}$ and water (40 ml). The reaction solution was heated on an oil bath for 5 h at 100°C, after which pyridine (10 ml) was added to the solution. Boiling H₂O (30 ml) was added and the solution was centrifuged. Green crystals of the title compound crystallized from the filtrate after several days.

S3. Refinement

C-bound H atoms were positioned geometrically and refined as riding atoms, with C—H = 0.97 Å and $U_{iso}(H) = 1.2U_{eq}(C)$. The H atoms attached to hydroxy groups and water molecules were located on a difference Fourier map and refined isotropically except H3, which was refined as riding, with O3—H3 = 0.82 Å and $U_{iso}(H3) = 1.5U_{eq}(O3)$. A 50% positional disorder was assigned to N1 and C4, which share a position of the pyridine ring, as this provided the best fit of the data. Short C—C bond interactions, probably due to this disorder, are observed for the pyridinium cation.



Figure 1

Molecular structure of the title compound. Displacement ellipsoids are drawn at the 50% probability level. [Symmetry code: (i) 1-x, 1-y, 1-z.]

Pyridinium diaquabis(methylenediphosphonato- $\kappa^2 O, O'$)chromate(III) tetrahydrate

Crystal data

$(C_{5}H_{6}N)[Cr(CH_{4}O_{6}P_{2})_{2}(H_{2}O)_{2}]\cdot 4H_{2}O$ $M_{r} = 588.17$ Triclinic, <i>P</i> 1 Hall symbol: -P 1 a = 7.206 (5) Å b = 7.485 (5) Å c = 10.984 (5) Å a = 107.085 (5)° $\beta = 106.128$ (5)° $\gamma = 94.496$ (5)° V = 535.7 (6) Å ³	Z = 1 F(000) = 303 $D_x = 1.823 \text{ Mg m}^{-3}$ Mo Ka radiation, $\lambda = 0.71073 \text{ Å}$ Cell parameters from 6300 reflections $\theta = 0.8 - 0.9^{\circ}$ $\mu = 0.92 \text{ mm}^{-1}$ T = 100 K Cuboid, green $0.22 \times 0.16 \times 0.08 \text{ mm}$
Data collection	
Bruker APEXII CCD diffractometer φ and ω scans Absorption correction: multi-scan (<i>SADABS</i> ; (Bruker, 2001) $T_{min} = 0.843, T_{max} = 0.931$ 8784 measured reflections	2632 independent reflections 2483 reflections with $I > 2\sigma(I)$ $R_{int} = 0.020$ $\theta_{max} = 28.3^{\circ}, \theta_{min} = 4.1^{\circ}$ $h = -9 \rightarrow 9$ $k = -9 \rightarrow 6$ $l = -14 \rightarrow 14$
Refinement Refinement on F^2 Least-squares matrix: full $R[F^2 > 2\sigma(F^2)] = 0.024$ $wR(F^2) = 0.070$ S = 1.05 2632 reflections	179 parameters 16 restraints H atoms treated by a mixture of independent and constrained refinement $w = 1/[\sigma^2(F_o^2) + (0.0342P)^2 + 0.4874P]$ where $P = (F_o^2 + 2F_c^2)/3$

 $(\Delta/\sigma)_{\rm max} < 0.001$ $\Delta \rho_{\rm max} = 0.47 \text{ e} \text{ Å}^{-3}$ $\Delta \rho_{\rm min} = -0.62 \text{ e } \text{\AA}^{-3}$

	x	У	Ζ	$U_{ m iso}$ */ $U_{ m eq}$	Occ. (<1)
Cr1	0.5000	0.5000	0.5000	0.0082 (3)	
P2	0.2173 (2)	0.79359 (19)	0.41325 (14)	0.0086 (4)	
P1	0.29362 (19)	0.7622 (2)	0.69142 (14)	0.0085 (4)	
O2	0.4491 (6)	0.6457 (6)	0.6635 (4)	0.0110 (8)	
O5	-0.0069 (6)	0.7069 (6)	0.3539 (4)	0.0128 (8)	
O3	0.3700 (6)	0.8943 (6)	0.8418 (4)	0.0134 (8)	
Н3	0.4774	0.9569	0.8570	0.020*	
O7	0.3373 (6)	0.6353 (6)	0.3947 (4)	0.0113 (8)	
O9	-0.0373 (8)	0.7358 (7)	0.8942 (5)	0.0235 (10)	
O1	0.7325 (6)	0.6953 (6)	0.5401 (4)	0.0121 (8)	
C1	0.2683 (8)	0.9145 (8)	0.5903 (6)	0.0108 (10)	
H4	0.3885	1.0063	0.6226	0.013*	
Н5	0.1633	0.9841	0.6027	0.013*	
O4	0.0957 (6)	0.6494 (6)	0.6673 (4)	0.0118 (8)	
O8	0.2601 (7)	0.9720 (7)	1.1133 (5)	0.0198 (10)	
N1	0.381 (7)	0.479 (8)	0.070 (5)	0.028 (12)	0.50
H1	0.3025	0.4691	0.1145	0.034*	0.50
C4	0.381 (8)	0.466 (10)	0.078 (5)	0.024 (10)	0.50
H4A	0.3051	0.4405	0.1294	0.029*	0.50
C3	0.5612 (13)	0.5896 (11)	0.1367 (8)	0.0312 (17)	
H3A	0.6025	0.6454	0.2295	0.037*	
C2	0.3206 (12)	0.3837 (11)	-0.0627 (8)	0.0322 (17)	
H2	0.2008	0.3014	-0.1034	0.039*	
O6	0.2543 (6)	0.9385 (6)	0.3492 (4)	0.0133 (8)	
H8	0.193 (16)	0.882 (13)	1.042 (9)	0.06 (3)*	
H7	0.262 (14)	0.950 (13)	1.184 (7)	0.03 (2)*	
H1B	0.844 (10)	0.677 (12)	0.577 (8)	0.03 (2)*	
H1A	0.731 (13)	0.806 (9)	0.573 (8)	0.025*	
H6	-0.032 (11)	0.597 (8)	0.352 (9)	0.03 (3)*	
H9A	0.002 (13)	0.710 (13)	0.828 (8)	0.03 (2)*	
H10	-0.110 (17)	0.817 (16)	0.888 (14)	0.07 (4)*	

Fractional atomic coordinates and isotropic or equivalent isotropic displacement parameters (\hat{A}^2)

Atomic displacement parameters $(Å^2)$

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
Cr1	0.0067 (6)	0.0081 (6)	0.0107 (6)	0.0020 (4)	0.0035 (4)	0.0035 (5)
P2	0.0072 (6)	0.0076 (7)	0.0120 (7)	0.0015 (5)	0.0031 (5)	0.0042 (5)
P1	0.0068 (6)	0.0084 (7)	0.0103 (7)	0.0011 (5)	0.0032 (5)	0.0025 (5)
O2	0.0103 (18)	0.0121 (19)	0.0119 (19)	0.0050 (15)	0.0043 (15)	0.0042 (15)
O5	0.0077 (18)	0.011 (2)	0.019 (2)	0.0004 (15)	0.0019 (15)	0.0053 (16)
O3	0.0110 (18)	0.013 (2)	0.0126 (19)	-0.0004 (15)	0.0036 (15)	0.0006 (16)
O7	0.0103 (18)	0.0116 (19)	0.0136 (19)	0.0046 (15)	0.0047 (15)	0.0048 (15)

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09	0.029 (3)	0.024 (3)	0.021 (2)	0.004 (2)	0.014 (2)	0.007 (2)	
01	0.0082 (18)	0.0088 (19)	0.019 (2)	0.0013 (15)	0.0034 (16)	0.0043 (16)	
C1	0.010(2)	0.008 (2)	0.013 (3)	0.0014 (19)	0.003 (2)	0.003 (2)	
O4	0.0085 (18)	0.0114 (19)	0.0153 (19)	0.0000 (14)	0.0039 (15)	0.0043 (15)	
08	0.018 (2)	0.026 (2)	0.013 (2)	-0.0028 (18)	0.0032 (17)	0.0052 (19)	
N1	0.04 (2)	0.028 (18)	0.033 (19)	0.019 (13)	0.019 (16)	0.023 (13)	
C4	0.026 (18)	0.019 (15)	0.018 (15)	-0.001 (12)	-0.008 (12)	0.008 (13)	
C3	0.042 (5)	0.020 (3)	0.021 (3)	0.007 (3)	-0.004 (3)	0.006 (3)	
C2	0.033 (4)	0.023 (4)	0.029 (4)	0.000 (3)	-0.008 (3)	0.009 (3)	
O6	0.0148 (19)	0.0108 (19)	0.016 (2)	0.0016 (15)	0.0053 (16)	0.0073 (16)	

Geometric parameters (Å, °)

Crl—Ol	1.991 (4)	O1—H1B	0.83 (6)	
Cr1—02	1.956 (4)	O1—H1A	0.80 (6)	
Cr1—O7	1.964 (4)	C1—H4	0.9700	
P2—O6	1.499 (4)	C1—H5	0.9700	
P2—O7	1.519 (4)	O8—H8	0.85 (7)	
P2—O5	1.568 (4)	O8—H7	0.83 (6)	
P2—C1	1.804 (6)	N1—C2	1.34 (5)	
P104	1.512 (4)	N1—C3	1.36 (4)	
P104	1.512 (4)	N1—H1	0.8600	
P1	1.515 (4)	C4—C3	1.40 (5)	
P1O3	1.568 (4)	C4—C2	1.41 (5)	
P1	1.797 (6)	C4—H4A	0.9300	
O5—H6	0.83 (5)	C3—C2 ⁱ	1.371 (13)	
О3—Н3	0.8200	С3—НЗА	0.9300	
О9—Н9А	0.83 (6)	C2—C3 ⁱ	1.371 (13)	
O9—H10	0.84 (7)	C2—H2	0.9300	
O2 ⁱⁱ —Cr1—O2	180.00 (15)	Р2—О5—Н6	114 (5)	
O2 ⁱⁱ —Cr1—O7	88.35 (17)	P1—O3—H3	109.5	
O2-Cr1-O7	91.65 (17)	P2—O7—Cr1	140.0 (3)	
O2 ⁱⁱ —Cr1—O7 ⁱⁱ	91.65 (17)	H9A—O9—H10	108 (10)	
O2-Cr1-O7 ⁱⁱ	88.35 (17)	Cr1—O1—H1B	119 (6)	
O7—Cr1—O7 ⁱⁱ	180.0 (2)	Cr1—O1—H1A	120 (6)	
O2 ⁱⁱ —Cr1—O1 ⁱⁱ	90.51 (17)	H1B—O1—H1A	107 (9)	
O2-Cr1-O1 ⁱⁱ	89.49 (17)	P1—C1—P2	114.8 (3)	
O7—Cr1—O1 ⁱⁱ	90.81 (18)	P1	108.6	
O7 ⁱⁱ —Cr1—O1 ⁱⁱ	89.19 (18)	P2	108.6	
O2 ⁱⁱ —Cr1—O1	89.49 (17)	P1—C1—H5	108.6	
02—Cr1—O1	90.51 (17)	P2—C1—H5	108.6	
07—Cr1—O1	89.19 (18)	H4—C1—H5	107.5	
O7 ⁱⁱ —Cr1—O1	90.81 (18)	H8—O8—H7	114 (10)	
Ol ⁱⁱ —Crl—Ol	180.0 (3)	C2—N1—C3	123 (4)	
O6—P2—O7	114.8 (2)	C2—N1—H1	118.3	
O6—P2—O5	107.8 (2)	C3—N1—H1	118.3	
O7—P2—O5	109.7 (2)	C3—C4—C2	116 (5)	

O6—P2—C1	108.2 (3)	C3—C4—H4A	122.1
O7—P2—C1	109.0 (2)	C2—C4—H4A	122.1
O5—P2—C1	107.1 (2)	$N1 - C3 - C2^{i}$	118 (2)
O4—P1—O2	115.5 (2)	$C2^{i}$ —C3—C4	122 (3)
O4—P1—O2	115.5 (2)	N1—C3—H3A	121.1
O4—P1—O3	107.9 (2)	C2 ⁱ —C3—H3A	121.1
O4—P1—O3	107.9 (2)	C4—C3—H3A	116.7
O2—P1—O3	108.6 (2)	$N1-C2-C3^{i}$	119 (2)
O4—P1—C1	110.1 (2)	$C3^{i}$ — $C2$ — $C4$	122 (2)
O4—P1—C1	110.1 (2)	N1—C2—H2	120.7
O2—P1—C1	107.6 (2)	C3 ⁱ —C2—H2	120.7
O3—P1—C1	106.9 (3)	C4—C2—H2	117.1
P1—O2—Cr1	134.1 (2)		

Symmetry codes: (i) -*x*+1, -*y*+1, -*z*; (ii) -*x*+1, -*y*+1, -*z*+1.

Hydrogen-bond geometry (Å, °)

<i>D</i> —H··· <i>A</i>	D—H	Н…А	D····A	D—H···A
С1—Н4…Об ^{ііі}	0.97	2.49	3.346 (7)	147
C4—H4A····O9 ^{iv}	0.93	2.16	2.93 (7)	140
N1—H1····O9 ^{iv}	0.86	2.32	3.03 (5)	141
O1—H1A···O6 ⁱⁱⁱ	0.80 (6)	1.83 (6)	2.634 (6)	176 (9)
O1— $H1B$ ···O4 ^v	0.83 (6)	1.87 (6)	2.704 (6)	177 (9)
O3—H3…O8 ^{vi}	0.82	1.83	2.629 (6)	163
O5—H6····O4 ^{iv}	0.83 (5)	1.80 (5)	2.619 (6)	175 (10)
O8—H7····O6 ^{vii}	0.83 (6)	1.86 (6)	2.687 (6)	171 (9)
O8—H8…O9	0.85 (7)	1.94 (8)	2.748 (7)	158 (11)
O9—H9A…O4	0.83 (6)	2.00 (6)	2.833 (6)	179 (10)
O9—H10…O8 ^{viii}	0.84 (7)	1.99 (7)	2.820 (7)	174 (13)

Symmetry codes: (iii) -x+1, -y+2, -z+1; (iv) -x, -y+1, -z+1; (v) x+1, y, z; (vi) -x+1, -y+2, -z+2; (vii) x, y, z+1; (viii) -x, -y+2, -z+2.