metal-organic papers

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Key indicators

Single-crystal X-ray study T = 298 KMean $\sigma(C-C) = 0.011 \text{ Å}$ R factor = 0.025 wR factor = 0.078 Data-to-parameter ratio = 15.9

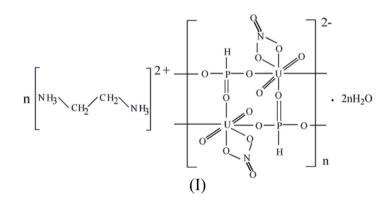
For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e.

catena-Poly[1,2-ethylenediammonium [[nitratouranyl]-µ₃-phosphito] dihydrate]

The crystal structure of the title compound, $\{(C_2H_{10}N_2)[U(H-PO_3)(NO_3)O_2]\cdot 2H_2O\}_n$, consists of polymeric uranyl complex anions, ethylenediammonium cations and uncoordinated water molecules. The polymeric uranyl complex anion displays a ladder-like structure, each UO₂ unit being coordinated by three phosphite dianions and one nitrate anion with a pentagonal-bipyramidal geometry. The ethylenediammonium cation is located on an inversion center and is hydrogen bonded with the uranyl complex.

Comment

Many uranyl phosphates have been reported because of their applications in fields such as ion exchange and protonic conduction (Morosin, 1978). However, only four uranyl phosphites have been reported in the Cambridge Structural Database (Allen, 2002; update of November 2005). We report here the structure of the title polymeric uranyl complex, (I), which is bridged by the phosphite dianion.



The crystal structure of (I) consists of polymeric uranyl complex anions, ethylenediammonium counter-cations and uncoordinated water molecules. The polymeric uranyl complex displays a ladder-like structure (Fig. 1). Each UO_2 unit is coordinated by three phosphite dianions and one nitrate anion and has a pentagonal-bipyramidal coordination geometry. The nitrate anion chelates the U^{VI} atom with longer U-O bond distances (Table 1). Each phosphite dianion bridges three UO₂ units, forming the polymeric complex chain. This is structurally similar to uranyl sulfate (Doran *et al.*, 2003) and uranyl selenate (Krivovichev & Kahlenberg, 2005).

The ethylenediammonium cation is located on an inversion center and is hydrogen bonded with the uranyl complex. The hydrogen bonds (Table 2) between the uncoordinated water molecule and the complex, and between the uncoordinated water and ethylenediammonium stabilize the crystal structure.

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Experimental

 $UO_2(NO_3)_2 \cdot 6H_2O$ (0.086 g, 0.17 mmol), H_3PO_3 (0.077 g, 0.94 mmol) and 1,2-ethylenediamine (0.041 g, 0.68 mmol) were mixed with distilled water (2 ml). The mixture was stirred at room temperature until completely homogenized. The solution was poured on to a watch glass and left to evaporate. Single crystals of (I) were obtained the next day.

Z = 1

 $D_x = 3.250 \text{ Mg m}^{-3}$

Cell parameters from 3075

 $0.58 \times 0.17 \times 0.06 \text{ mm}$

2086 independent reflections

1949 reflections with $I > 2\sigma(I)$

Mo $K\alpha$ radiation

reflections $\theta = 2.3-27.5^{\circ}$ $\mu = 17.43 \text{ mm}^{-1}$

T = 298 (2) K Needle, yellow

 $\begin{aligned} R_{\text{int}} &= 0.038\\ \theta_{\text{max}} &= 27.5^{\circ}\\ h &= -9 \rightarrow 8\\ k &= -9 \rightarrow 9 \end{aligned}$

 $l = -12 \rightarrow 11$

Crystal data

$(C_2H_{10}N_2)[U(HPO_3)(NO_3)O_2]$
2H ₂ O
$M_r = 922.18$
Triclinic, P1
a = 7.0106 (17) Å
b = 7.643 (2) Å
c = 9.846 (1) Å
$\alpha = 108.522 (7)^{\circ}$
$\beta = 100.469 \ (8)^{\circ}$
$\gamma = 102.125 \ (9)^{\circ}$
V = 471.20 (18) Å ³

Data collection

Rigaku R-AXIS RAPID
diffractometer
ω scans
Absorption correction: multi-scan
(ABSCOR; Higashi, 1995)
$T_{\min} = 0.040, \ T_{\max} = 0.342$
3928 measured reflections

Refinement

Refinement on F^2	$w = 1/[\sigma^2(F_0^2) + (0.0529P)^2]$
$R[F^2 > 2\sigma(F^2)] = 0.025$	+ 0.6522P]
$wR(F^2) = 0.078$	where $P = (F_0^2 + 2F_c^2)/3$
S = 1.17	$(\Delta/\sigma)_{\rm max} = 0.002$
2086 reflections	$\Delta \rho_{\rm max} = 1.53 \ {\rm e} \ {\rm \AA}^{-3}$
131 parameters	$\Delta \rho_{\rm min} = -1.31 \text{ e } \text{\AA}^{-3}$
H-atom parameters constrained	

Table 1

Selected bond lengths (Å).

U1-01	1.784 (5)	$U1 - O5^{ii}$	2.307 (5)
U1-O2	1.772 (6)	U1-O6	2.534 (6)
U1-O3	2.313 (5)	U1-07	2.526 (5)
$U1-O4^{i}$	2.298 (5)		

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

Table 2			
Hydrogen-bond	geometry	′ (Å,	°).

$D - H \cdots A$	D-H	$H \cdot \cdot \cdot A$	$D \cdot \cdot \cdot A$	$D - \mathbf{H} \cdot \cdot \cdot A$
$N2-H2A\cdots OW1^{iii}$	0.91	2.22	2.955 (10)	138
$N2-H2A\cdots OW1^{iv}$	0.91	2.30	2.893 (10)	123
$N2-H2B\cdots O6^{ii}$	0.91	2.33	3.104 (10)	143
$N2-H2C\cdots O3^{v}$	0.91	2.49	3.136 (9)	128
$N2-H2C\cdots O5^{v}$	0.91	2.10	2.975 (9)	161
$OW1-H1W \cdot \cdot \cdot O4^{vi}$	0.82	2.37	3.142 (9)	157
$OW1-H2W \cdot \cdot \cdot O2^{vi}$	0.82	2.31	3.080 (9)	156

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

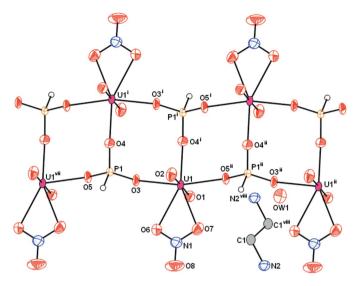


Figure 1

The molecular structure of (I), shown with 50% probability displacement ellipsoids. H atoms on the cation and on the uncoordinated water molecule have been omitted for clarity. [Symmetry codes: (i) -x, -y + 1, -z; (ii) x + 1, y, z; (vii) x - 1, y, z; (viii) -x + 2, -y + 2, -z + 1.]

H atoms on the water molecule and phosphite dianion were located in a difference Fourier map and refined as riding in their asfound relative positions, with $U_{iso}(H) = 1.2U_{eq}(O,P)$. Methyl H atoms were placed in calculated positions, and the torsion angles refined to fit the electron density, with $U_{iso}(H) = 1.5U_{eq}(C)$. Methylene H atoms were placed in calculated positions and refined as riding, with $U_{iso}(H) = 1.2U_{eq}(C)$. The highest peak is 0.87 Å from atom U1 and the deepest hole is 1.27 Å from atom O5.

Data collection: *PROCESS-AUTO* (Rigaku, 1998); cell refinement: *PROCESS-AUTO*; data reduction: *CrystalStructure* (Rigaku/ MSC, 2002); program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *SHELXTL* (Sheldrick, 2001); software used to prepare material for publication: *SHELXTL*.

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