inorganic papers

Acta Crystallographica Section E Structure Reports Online

ISSN 1600-5368

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Key indicators

Single-crystal X-ray study T = 296 KMean $\sigma(\text{Na-O}) = 0.015 \text{ Å}$ R factor = 0.056 wR factor = 0.098 Data-to-parameter ratio = 14.3

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

Poly[diammonium diaquahexahydroxyoctadecaoxodisodiohexamolybdonickelate(II)]

The title compound, $(NH_4)_2[Na_2Ni(OH)_6Mo_6O_{18}(H_2O)_2]$, features inversion-generated hexamolybdonickelate(II) anions (Ni site symmetry 1). The anions are linked by the five-coordinate sodium cations into layers. Adjacent layers are linked through $N-H\cdots O$ and $O-H\cdots O$ hydrogen bonds involving the ammonium ions and water molecules, thus forming a three-dimensional network.

Comment

Varying the inorganic or organic cation employed in the synthesis is an efficient strategy for producing novel polyoxometalates (Liu, Li *et al.*, 2006). To extend our recent work, where hexamolybdonickelate(II) units are linked by cations forming three-dimensional networks (Liu, Zhou *et al.*, 2006), we obtained the title compound, (I), in which the anions of these compounds are isostructural.

The title compound, (I), is composed of $[Ni(OH)_6Mo_6O_{18}]^{4-}$ anionic clusters, sodium and ammonium cations, and water molecules as shown in Fig. 1. The $[Ni(OH)_6Mo_6O_{18}]^{4-}$ cluster adopts a β -type Anderson structure, made up of seven edge-sharing octahedra: six MoO₆ groups arranged hexagonally around the central Ni(OH)₆ octahedron (Ni site symmetry $\overline{1}$). The Ni-O and Mo-O distances vary between 2.015 (7)-2.023 (7) Å and 1.659 (7)-2.250 (7) Å, respectively (Table 1). Bond valence sum calculations (Brown & Altermatt, 1985) indicated oxidation states of 5.95-6.07 for the Mo atoms and 1.97 for Ni in good agreement with the expected values of 6 and 2, respectively.



Figure 1

Fragment of (I) showing the polyoxometallate cluster and one attached NaO₅ grouping. Displacement ellipsoids are drawn at the 30% probability level (H atoms omitted for clarity). [Symmetry code: (i) 1 - x, -y, -z.]

© 2007 International Union of Crystallography All rights reserved Received 6 December 2006 Accepted 11 December 2006 The Na⁺ cation is bonded to three terminal O atoms from two neighboring $[Ni(OH)_6Mo_6O_{18}]^{4-}$ polyanions and two water molecules, resulting in an NaO₅ square-based pyramid. Thus, the polyanions and Na⁺ cations form a 4,4-connected layered structure. These are further linked through ammonium ions and water molecules by hydrogen bonds, thus forming a three-dimensional framework (Fig. 2).

Experimental

 $(NH_4)_6Mo_7O_{24}$ ·6H₂O (2 g, 0.16 mmol) and NiSO₄·6H₂O (0.050 g, 0.2 mmol) were dissolved in water (25 ml) with stirring. NaCl (0.10 g, 1.7 mmol) was then added to the transparent solution and the pH was adjusted to about 2.5–2.6 by adding 4 *M* HCl. The mixture was refluxed at 343 K for about 5 h and then cooled to room temperature; the solution was filtered into a 50 ml beaker. Slow evaporation of the solvent at room temperature led to green block-shaped crystals of (I) suitable for X-ray diffraction after two weeks.

Crystal data		Та
$(NH_4)_2[Na_{2-}]$	V = 1361.1 (2) Å ³	H
$Ni(OH)_6Mo_6O_{18}(H_2O)_2$	Z = 2	
$M_r = 1214.54$	$D_{\rm r} = 2.964 {\rm Mg} {\rm m}^{-3}$	D ·
Monoclinic, $P2_1/c$	Mo $K\alpha$ radiation	
a = 11.7792 (10) Å	$\mu = 3.50 \text{ mm}^{-1}$	N-
b = 10.9353 (9) Å	T = 296 (2) K	N-
c = 11.7793 (10) Å	Block light green	N-
$\beta = 116.223 (A)^{\circ}$	$0.16 \times 0.12 \times 0.11 \text{ mm}$	N -
p = 110.225 (4)	$0.10 \times 0.12 \times 0.11$ mm	01
		02
Data collection		03
Bruker SMADT ADEX II CCD	7181 measured reflections	O
DIUKEI SMART ALLA II CCD	/181 measured reflections	01

2674 independent reflections

 $R_{\rm int} = 0.036$

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

1396 reflections with $I > 2\sigma(I)$

diffractometer ω scans Absorption correction: multi-scan (*SADABS*; Bruker, 2002) $T_{min} = 0.605, T_{max} = 0.680$

Refinement

Refinement on F^2	H-atom parameters constrained
$R[F^2 > 2\sigma(F^2)] = 0.056$	$w = 1/[\sigma^2(F_0^2) + (0.038P)^2]$
$wR(F^2) = 0.098$	where $P = (F_0^2 + 2F_c^2)/3$
S = 0.95	$(\Delta/\sigma)_{\rm max} < 0.001$
2674 reflections	$\Delta \rho_{\rm max} = 0.75 \ {\rm e} \ {\rm \AA}^{-3}$
187 parameters	$\Delta \rho_{\rm min} = -0.67 \ {\rm e} \ {\rm \AA}^{-3}$

Та	ble	e 1	l
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Selected bond lengths (Å).

Ni-O1	2.015 (7)	Mo2-O10	1.922 (7)
Ni-O3	2.022 (7)	Mo2-O3	2.204 (7)
Ni-O2	2.023 (7)	Mo2-O1 ⁱ	2.250 (7)
Ni-O1 ⁱ	2.015 (7)	Mo3-O8	1.674 (7)
Ni-O3 ⁱ	2.022 (7)	Mo3-O9	1.702 (7)
Ni-O2 ⁱ	2.023 (7)	Mo3-O10	1.934 (7)
Mo1-O6	1.659 (7)	Mo3-O7	1.935 (7)
Mo1-O5	1.698 (8)	Mo3-O2	2.193 (7)
Mo1-O4	1.933 (7)	Mo3-O3	2.211 (7)
Mo1-O7	1.937 (7)	Na-OW3	2.045 (14)
Mo1-O2	2.196 (7)	Na-OW1	2.258 (16)
Mo1-O1	2.209 (7)	Na-O5	2.350 (13)
Mo2-O11	1.684 (8)	Na-O12 ⁱ	2.389 (13)
Mo2-O12	1.712 (8)	Na-O11 ⁱⁱ	2.442 (13)
$M_02 - O4^i$	1 909 (8)		. ,

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



Figure 2 The packing for (I), with hydrogen bonds indicated by dashed lines.

Table 2	
Hydrogen-bond geometry (Å, $^{\circ}$).	

$D - H \cdots A$	$D-{\rm H}$	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdot \cdot \cdot A$
N-HN1···O10	0.84	1.88	2.681 (12)	157
N-HN2···O9 ⁱⁱⁱ	0.93	2.03	2.830 (12)	143
$N-HN3\cdots O5^{iv}$	0.84	2.06	2.883 (12)	164
$N-HN4\cdots O12^{v}$	0.84	2.01	2.826 (12)	164
$O1-H1\cdots O6^{vi}$	0.87	1.98	2.847 (10)	169
$O2-H2\cdots OW2$	0.88	1.82	2.649 (11)	156
$O3-H3\cdots O8^{vi}$	0.87	1.91	2.781 (11)	171
$OW1 - HW1B \cdot \cdot \cdot O10^{vii}$	0.85	2.34	3.074 (14)	146
$OW2-HW2B\cdots O7^{iv}$	0.91	2.18	2.864 (10)	131
OW3−HW3A···O9 ^{viii}	0.86	2.36	2.891 (13)	121
$OW3-HW3B\cdots O4^{vi}$	0.84	2.13	2.690 (12)	124

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

All the H atoms were located in difference maps and refined in their as-found relative positions (O-H = 0.82-0.91 Å and N-H = 0.84-0.93 Å), using a riding model with U_{iso} (H) = 1.2 U_{eq} (O,N).

Data collection: *SMART* (Bruker, 2002); cell refinement: *SAINT* (Bruker, 2002); data reduction: *SAINT*; program(s) used to solve structure: *SHELXTL* (Bruker, 1997); program(s) used to refine structure: *SHELXTL*; molecular graphics: *DIAMOND* (Brandenburg, 1999); software used to prepare material for publication: *SHELXTL*.

This work was supported by Jilin Normal University Innovative Foundation (research grant No. 20050051006).

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