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Key indicators

Single-crystal X-ray study
 $T = 298\text{ K}$
Mean $\sigma(\text{I-O}) = 0.004\text{ \AA}$
 R factor = 0.037
 wR factor = 0.087
Data-to-parameter ratio = 17.2For details of how these key indicators were
automatically derived from the article, see
<http://journals.iucr.org/e>.Sodium barium hexahydrogenhexamolybdo-
aluminate(III) decahydrate

The title compound, $\text{NaBa}[\text{AlMo}_6(\text{OH})_6\text{O}_{18}] \cdot 10\text{H}_2\text{O}$, contains a B-type Anderson polyanion $[\text{AlMo}_6(\text{OH})_6\text{O}_{18}]^{3-}$ having an inversion center with approximate D_{3d} symmetry. There are two uncoordinated water molecules per formula unit, the remaining eight being coordinated to the Na^+ and Ba^{2+} cations, which link the polyanions to form the three-dimensional structure. Hydrogen-bonding interactions exist to stabilize the structure.

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Comment

The well known Anderson-type polyanions (Anderson, 1937) can be found in many compounds employing rare-earth cations and transition-metal cations or complexes as bridging linkers (Gao *et al.*, 2006; Shivaiah *et al.*, 2003). In contrast, the use of alkaline-earth cations as linkers remains rare, as in the title compound, $\text{NaBa}[\text{AlMo}_6(\text{OH})_6\text{O}_{18}] \cdot 10\text{H}_2\text{O}$ (I).

The structure consists of an Anderson polyanion, a sodium ion, a barium ion, eight coordinated waters, and two uncoordinated waters. The $[\text{AlMo}_6(\text{OH})_6\text{O}_{18}]^{3-}$ polyanion adopts a B-type Anderson structure, consisting of seven edge-sharing octahedra, six of which are $\{\text{MoO}_6\}$ octahedra arranged hexagonally around the central $\{\text{Al}(\text{OH})_6\}$ octahedron (Fig. 1). It has an inversion center and is close to D_{3d} symmetry, all bond lengths and angles being within the normal ranges (Lee *et al.*, 1991; Liu *et al.*, 2006). The Na^+ ion is six-coordinate and the Ba^{2+} ion is eight-coordinate. Not only do these cations

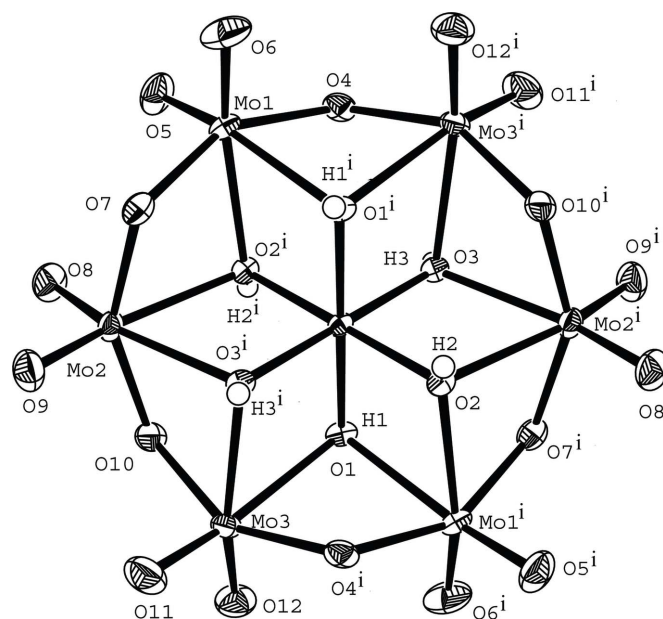


Figure 1
Structure of the polyanion in (I), showing 50% probability displacement ellipsoids. [Symmetry code: (i) $-x + \frac{3}{2}, -y + \frac{3}{2}, -z$.]

compensate for the polyanionic charge, they also act as linkers to form the three-dimensional structure (Fig. 2). It is believed that extensive hydrogen bonding interactions play an important role in stabilizing the structure.

Experimental

To a 30 ml aqueous solution of $\text{AlCl}_3 \cdot 6\text{H}_2\text{O}$ (0.36 g, 1.5 mmol) were added 10 ml aqueous solution of $\text{Na}_2\text{MoO}_4 \cdot 2\text{H}_2\text{O}$ (0.90 g, 3.7 mmol) and 5 ml of glacial acetic acid, followed by addition of 10 ml $\text{BaCl}_2 \cdot 2\text{H}_2\text{O}$ (0.15 g, 0.6 mmol). The pH of the resulting mixture was adjusted with dilute hydrochloric acid to about 2.6 and it was stirred for half an hour. The filtrate was allowed to evaporate at room temperature. After five days, colorless block-shaped crystals of (I) were collected in about 64% yield (0.53 g, based on Mo).

Crystal data

$\text{NaBa}[\text{AlMo}_6(\text{OH})_6\text{O}_{18}] \cdot 10\text{H}_2\text{O}$	$V = 3037.0 (10) \text{ \AA}^3$
$M_r = 1333.16$	$Z = 4$
Monoclinic, $C2/c$	Mo $K\alpha$ radiation
$a = 23.553 (5) \text{ \AA}$	$\mu = 3.84 \text{ mm}^{-1}$
$b = 11.549 (2) \text{ \AA}$	$T = 298 (2) \text{ K}$
$c = 11.227 (2) \text{ \AA}$	$0.14 \times 0.13 \times 0.12 \text{ mm}$
$\beta = 96.06 (3)^\circ$	

Data collection

Rigaku R-AXIS RAPID IP diffractometer	14015 measured reflections
Absorption correction: multi-scan (ABSCOR; Higashi, 1995)	3398 independent reflections
$T_{\min} = 0.588$, $T_{\max} = 0.632$	3082 reflections with $I > 2\sigma(I)$
	$R_{\text{int}} = 0.045$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.037$	1 restraint
$wR(F^2) = 0.087$	H-atom parameters constrained
$S = 1.14$	$\Delta\rho_{\text{max}} = 1.75 \text{ e \AA}^{-3}$
3398 reflections	$\Delta\rho_{\text{min}} = -0.89 \text{ e \AA}^{-3}$
198 parameters	

Table 1

Selected geometric parameters (\AA , $^\circ$).

Al1—O1	1.902 (4)	Mo2—O7	1.959 (4)
Al1—O2	1.894 (4)	Mo2—O8	1.705 (4)
Al1—O3	1.896 (4)	Mo2—O9	1.708 (4)
Mo1—O4	1.936 (4)	Mo3—O10	1.935 (4)
Mo1—O5	1.704 (4)	Mo3—O11	1.702 (5)
Mo1—O6	1.713 (5)	Mo3—O12	1.712 (5)
O2—Al1—O3	84.38 (16)	O8—Mo2—O10	98.7 (2)
O3—Al1—O1	94.92 (16)	O10—Mo2—O7	150.75 (17)
O5—Mo1—O4	97.4 (2)	O11—Mo3—O12	105.7 (2)
O7—Mo1—O4	150.57 (17)	O11—Mo3—O1	160.56 (19)

Table 2

Hydrogen-bond geometry (\AA , $^\circ$).

$D-H \cdots A$	$D-H$	$H \cdots A$	$D \cdots A$	$D-H \cdots A$
O1—H1 \cdots O8 ⁱ	0.85	1.95	2.790 (6)	169
O2—H2 \cdots O5W ⁱⁱ	0.85	1.83	2.615 (6)	153
O3—H3 \cdots O5 ⁱ	0.85	1.88	2.718 (6)	169
O1W—H4 \cdots O6 ⁱⁱⁱ	0.85	2.14	2.769 (7)	130
O1W—H5 \cdots O4W ^{iv}	0.85	1.93	2.772 (7)	169

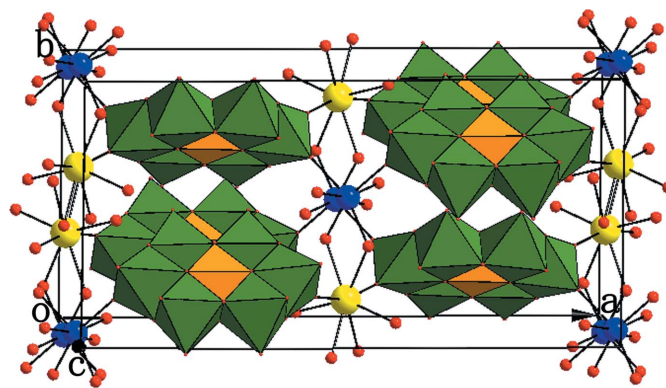


Figure 2

Polyhedral representation of the stacking of (I) in the unit cell. The uncoordinated waters and H atoms have been omitted for clarity. Blue circles are Na^+ and blue-gray circles are Ba^{2+} .

$D-H \cdots A$	$D-H$	$H \cdots A$	$D \cdots A$	$D-H \cdots A$
O2W—H6 \cdots O9 ^v	0.85	2.50	3.211 (7)	142
O2W—H7 \cdots O3W ^{vi}	0.85	2.06	2.895 (8)	165
O3W—H8 \cdots O9 ^{vii}	0.85	2.43	3.114 (7)	139
O3W—H9 \cdots O10 ^v	0.85	1.90	2.738 (7)	167
O4W—H10 \cdots O9 ^{viii}	0.85	2.05	2.848 (7)	157
O4W—H11 \cdots O4 ^{ix}	0.85	1.91	2.747 (6)	170
O5W—H12 \cdots O7 ^x	0.85	2.07	2.810 (7)	145
O5W—H13 \cdots O7 ^{xi}	0.85	2.20	3.003 (6)	157

Symmetry codes: (i) $x, -y + 1, z - \frac{1}{2}$; (ii) $-x + 1, -y + 2, -z$; (iii) $-x + \frac{3}{2}, y + \frac{1}{2}, -z + \frac{1}{2}$; (iv) $-x + 2, -y + 2, -z$; (v) $-x + 2, y + 1, -z + \frac{1}{2}$; (vi) $x, -y + 3, z - \frac{1}{2}$; (vii) $-x + 2, -y + 2, -z + 1$; (viii) $-x + 2, y, -z + \frac{1}{2}$; (ix) $x + \frac{1}{2}, y + \frac{1}{2}, z$; (x) $-x + 1, y, -z + \frac{1}{2}$; (xi) $x - \frac{1}{2}, -y + \frac{3}{2}, z - \frac{1}{2}$.

H atoms of the hydroxyl groups around the Al^{3+} ion and on water molecules were located in difference Fourier maps, idealized with $\text{O—H} = 0.85 \text{ \AA}$, and refined as riding, with $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{O})$.

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

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