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Key indicators

Single-crystal X-ray study T = 293 KMean $\sigma(C-C) = 0.012 \text{ Å}$ H-atom completeness 46% R factor = 0.041 wR factor = 0.112 Data-to-parameter ratio = 12.4

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

A polyoxomolybdate coordinated by glycine ligands: $K_4[Mo_8O_{26}(NH_3CH_2COO)_2] \cdot 6H_2O$

The title compound, tetrapotassium bis(glycine)hexacosaoxooctamolybdenum(VI) hexahydrate, $K_4[Mo_8O_{26}(NH_3CH_2-COO)_2]\cdot 6H_2O$, consists of a molecular network of centrosymmetric $[Mo_8O_{26}(NH_3CH_2COO)_2]^{4-}$ anions, K^+ cations and water molecules, which interact by an extensive hydrogen-bonding network. The $[Mo_8O_{26}(NH_3CH_2COO)_2]^{4-}$ anion is constructed from MoO_6 and $MoO_5(OOCCH_2NH_3)$ octahedral units, where the latter has O atoms of terminal and bridging oxo groups as well as those of glycine ligands. Received 12 May 2004 Accepted 1 June 2004 Online 12 June 2004

Comment

There is increasing interest in polyoxometalates, partly due to their interactions with enzymes and with different organic ligands, which enable a better understanding of the antitumor/ viral activities of this class of compounds and modeling substrates involved in enzyme inhibition (Rhule et al., 1998; Inoue & Yamase, 1995; Crans et al., 1994). Thus, the reactivities of amino acids with polyoxometalates, as model studies for polyoxometalate-protein interactions, have attracted increasing attention. Some examples of structurally characterized polyoxoanions with covalently bound amino acids, $[MMo_6O_{21}(L)_3]^{n}$ (M = V, P, Se, Te, As, Sb, Bi; L = glycine, β -4-aminobutyric alanine. acid, L-lysine), [Mo₈O₂₆- $(L-lysH_2)_2]^{2-}$, $[Mo_{154}O_{462}H_{14}(H_2O)_{48}(HO_2C-(NH^{3+})HC CH_2-S-S-CH_2-CH(NH^{3+})-COO-)_{11}]^{3-}$, have been reported recently (Kortz et al., 2002, 2003; Müller et al., 2001). We report here the synthesis and the crystal structure of $K_4[Mo_8O_{26}(NH_3CH_2COO)_2]\cdot 6H_2O$, (I) (Fig. 1), which contains octamolybdate coordinated by glycine.



In the centrosymmetric $[(NH_3CH_2COO)_2Mo_8O_{26}]^{4-}$ anion of the title compound, the β -octamolybdate is coordinated by two glycines, each via one O atom of the carboxyl group. Bond-valence sum calculations (Brown & Altermatt, 1985;

© 2004 International Union of Crystallography Printed in Great Britain – all rights reserved Brese & O'Keeffe, 1991) indicated the valences of Mo and O to be +6 and -2, respectively. To keep the compound electrically neutral, three H atoms must be linked to each N atom.

The packing of (I) (Fig. 2) shows a network of weak N– H···O–Mo hydrogen bonds, with N···O = 2.851 (9)– 3.064 (9) Å. In the crystal structure, the K1 cation is surrounded by nine O atoms, *viz.* eight from different polyoxoanions and one from a water molecule. The K2 cation is surrounded by eight O atoms, *viz.* four from different polyoxoanions and four from water molecules.



Figure 1

The molecular structure of (I) (50% displacement ellipsoids). H atoms have been omitted. The prime corresponds to symmetry code (i) in Table 1



Figure 2

Packing diagram of (I), showing the hydrogen bonds as dashed lines (15% displacement ellipsoids).

Experimental

A mixture of K_2MoO_4 (5 mmol), KCl (5 mmol), H_2NCH_2COOH (15 mmol), N_2H_4 ·2HCl (0.5 mmol) and H_2O (10 ml) was sealed in a Teflon-lined stainless steel reactor and heated at 413 K for 4 d. After the reaction system was slowly cooled down to room temperature, small yellow crystals of (I) were obtained.

Crystal data

 $K_4[Mo_8O_{26}(C_2H_5NO_2)_2]\cdot 6H_2O$ Z = 1 $M_r = 1598.16$ $D_x = 3.020 \text{ Mg m}^{-3}$ Triclinic, $P\overline{1}$ Mo $K\alpha$ radiation a = 8.1065 (3) Å Cell parameters from 4462 b = 9.8767 (2) Å reflections c = 12.4936(5) Å $\theta = 1.70 - 25.04$ $\mu = 3.35 \text{ mm}^{-1}$ $\alpha = 98.638(2)^{\circ}$ $\beta = 99.295(2)^{\circ}$ T = 293 (2) K $\gamma = 113.606 (1)^{\circ}$ Block, yellow $V = 878.66 (5) \text{ Å}^3$ $0.22 \times 0.14 \times 0.10 \text{ mm}$ Data collection 3026 independent reflections

Bruker SMART CCD diffractometer ω scans Absorption correction: multi-scan (*SADABS*; Sheldrick, 1996) $T_{min} = 0.523, T_{max} = 0.715$ 4462 measured reflections

Refinement

Refinement on F^2 $R[F^2 > 2\sigma(F^2)] = 0.041$ $wR(F^2) = 0.112$ S = 1.053026 reflections 245 parameters H-atom parameters constrained $\theta_{\text{max}} = 25.0^{\circ}$ $h = -9 \rightarrow 9$ $k = -11 \rightarrow 11$ $l = -13 \rightarrow 14$ $w = 1/[\sigma^{2}(F_{o}^{2}) + (0.0581P)^{2}$ + 0.0251Pl

2740 reflections with $I > 2\sigma(I)$

 $R_{\rm int} = 0.026$

+ 9.0351*P*] where $P = (F_o^2 + 2F_c^2)/3$ $(\Delta/\sigma)_{max} = 0.010$ $\Delta\rho_{max} = 0.80 \text{ e} \text{ Å}^{-3}$ $\Delta\rho_{min} = -1.03 \text{ e} \text{ Å}^{-3}$ Extinction correction: *SHELXL*97 Extinction coefficient: 0.0011 (4)

Table 1

Selected bond distances (Å).

Mo1-O5	1.709 (5)	Mo3-O11	1.697 (5)
Mo1-O6	1.717 (5)	Mo3-O12	1.741 (5)
Mo1-O4	1.933 (5)	Mo3-O3	1.900 (5)
Mo1-O1 ⁱ	1.975 (5)	Mo3-O2	1.941 (5)
Mo1-O3 ⁱ	2.251 (5)	Mo3-O7	2.154 (5)
Mo1-O2	2.264 (5)	Mo3-O2 ⁱ	2.431 (5)
Mo2-O10	1.711 (5)	Mo4-O13	1.706 (5)
Mo2-O9	1.720 (5)	Mo4-O14	1.711 (5)
Mo2-O1	1.910 (5)	Mo4-O7	1.912 (5)
Mo2-O8	2.075 (5)	Mo4-O4	1.931 (5)
Mo2-O7	2.106 (5)	Mo4-O2	2.251 (5)
Mo2-O3	2.222 (5)	Mo4-O12 ⁱ	2.341 (5)

Symmetry code: (i) -x, -y, -z.

Table 2	
Hydrogen-bonding geon	netry (Å, °).

$D - H \cdots A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - H \cdots A$
$V1 - H1A \cdots O6^{ii}$ $V1 - H1B \cdots O10^{iii}$ $V1 - H1C \cdots O14^{iv}$ $V1 - H1C \cdots O10^{iv}$	0.89 0.89 0.89 0.89	2.15 2.25 2.26 2.39	2.936 (9) 3.000 (9) 2.851 (9) 3.064 (9)	147 142 124 133

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

H atoms on C and N were positioned geometrically (N-H = 0.89)and C-H = 0.97 Å), and refined as riding, with $U_{iso} = 1.5U_{eq}(N)$ or $1.2U_{eq}(C)$. Water H atoms were not located. Data collection: *SMART* (Siemens, 1994); cell refinement: *SAINT* (Siemens, 1994); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS*97 (Sheldrick, 1997); program(s) used to refine structure: *SHELXL*97; molecular graphics: *SHELXL*97 ; software used to prepare material for publication: *SHELXL*97.

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