

## A three-dimensional supramolecular vanadium hydroxylamide complex: poly[di- $\mu_2$ -aqua-bis(hydroxylamido)- $\mu_3$ -malonato-oxidosodiumvanadium(V)]

Qi-Ying Zhang, Heng-Qiang Zhang, Ai-Guo Kong, Qian Yang and Yong-Kui Shan\*

Department of Chemistry, East China Normal University, Shanghai 200062, People's Republic of China

Correspondence e-mail: ykshan@chem.ecnu.edu.cn

Received 17 July 2009

Accepted 8 September 2009

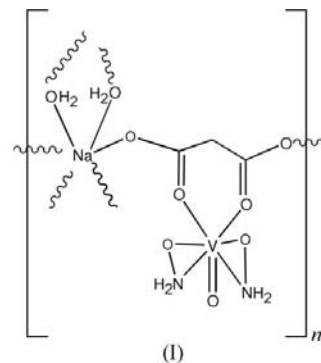
Online 26 September 2009

The crystal structure of the title compound,  $[\text{NaV}(\text{C}_3\text{H}_2\text{O}_4)(\text{NH}_2\text{O})_2\text{O}(\text{H}_2\text{O})_2]$ , is built up of  $\text{NaO}_6$  and  $\text{VO}_5\text{N}_2$  polyhedra connected through malonate bridges. The  $\text{NaO}_6$  octahedra are linked by edge sharing in the equatorial plane to form one-dimensional infinite chains. These chains are linked together by the malonate bridges to form two-dimensional layers. The distorted  $\text{VO}_5\text{N}_2$  pentagonal bipyramid is grafted on to the layer by a malonate carboxylate O atom. Adjacent layers are connected through  $\text{O}-\text{H}\cdots\text{O}$  and  $\text{N}-\text{H}\cdots\text{O}$  hydrogen bonds to build up a three-dimensional supramolecular structure.

### Comment

Simple vanadium salts such as  $\text{NaVO}_3$  and  $\text{VOSO}_4$  can lower blood glucose levels by activating glucose uptake by cells for metabolism in humans, but numerous studies have shown that organic vanadium complexes are less toxic and several times more effective in lowering blood glucose levels (Thompson *et al.*, 2002). Therefore, much research has gone into exploring the synthesis and structure of insulin-mimetic vanadium complexes. Vanadium hydroxylamide compounds are known to be promising candidates in the study of the insulin-mimetic activity of vanadium compounds (Tracey, 2000). Several vanadium hydroxylamide compounds have been reported, such as  $[\text{VO}(\text{NH}_2\text{O})(\text{dipic})(\text{H}_2\text{O})]$  (dipic is dipicolinic acid; Nuber *et al.*, 1981),  $[\text{VO}(\text{NH}_2\text{O})_2L]\cdot\text{H}_2\text{O}$  ( $L$  = glycine, serine and glycylglycine),  $[\text{VO}(\text{NH}_2\text{O})_2(\text{imidazole})]\text{Cl}$  (Keramidas *et al.*, 1997), but no vanadium hydroxylamide complexes with carboxylate ligands have been reported to date. Investigation of the preparation and crystal structure of vanadium hydroxylamide complexes with malonic acid provides not only useful information on vanadium chemistry but also promising new candidates for the study of the insulin-mimetic activity of vanadium. Therefore, we report here the preparation and

crystal structure of the title vanadium hydroxylamide complex with malonic acid, (I).



In the structure of (I), the  $\text{V}^{\text{V}}$  ion is seven-coordinated in a pentagonal bipyramidal geometry by two bidentate hydroxylamide ligands, one oxide ligand and two O atoms from a malonate ligand (Fig. 1). The hydroxylamide ligands coordinate in a side-on manner, as observed in related structures (Paul *et al.*, 1997; Keramidas *et al.*, 1997; Nuber *et al.*, 1981). The malonate behaves as a chelating ligand to the  $\text{V}^{\text{V}}$  ion. The centroids of the two hydroxylamide ligands and atom O2 of the malonate define the equatorial plane perpendicular to the  $\text{V}=\text{O}$  bond. The other chelating atom, O1, is in an axial position *trans* to the oxide ligand, with an axial  $\text{O1}-\text{V1}-\text{O7}$  angle of  $171.06(5)^\circ$ . The terminal  $\text{V1}=\text{O7}$  bond length is  $1.6017(10) \text{ \AA}$ , leading to the expected *trans* lengthening of the  $\text{V1}-\text{O1}$  bond to  $2.1470(9) \text{ \AA}$ , which is longer than the  $\text{V1}-\text{O2}$  bond (Table 1). The  $\text{O}-\text{N}$ ,  $\text{V}-\text{O}$  and  $\text{V}-\text{N}$  distances and  $\text{O}-\text{V}-\text{N}$  angles involving the hydroxylamide ligands are comparable, within experimental errors of  $0.0004 \text{ \AA}$  and

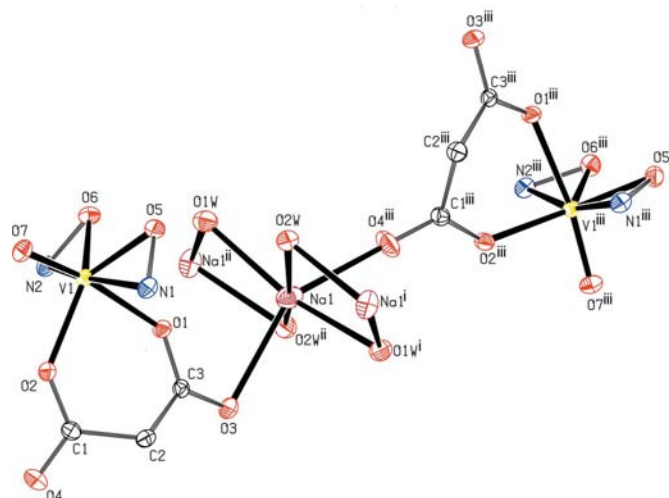


Figure 1

A view of part of the layer structure of (I), showing the asymmetric unit, atom connectivities and coordination environment. Displacement ellipsoids are drawn at the 30% probability level. H atoms have been omitted for clarity. [Symmetry codes: (i)  $x, -y + \frac{1}{2}, z - \frac{1}{2}$ ; (ii)  $x, -y + \frac{1}{2}, z + \frac{1}{2}$ ; (iii)  $x + 1, -y + \frac{1}{2}, z + \frac{1}{2}$ .]

0.0002°, respectively, with related vanadium hydroxylamide complexes reported in the literature (Table 3).

The coordination environment around the Na<sup>I</sup> ion can be described as a distorted octahedron. The vertices are occupied by six O atoms, of which four belong to water molecules [Na—O = 2.4132 (13)–2.4305 (13) Å] located in the equatorial plane, with two slightly more distant O atoms from two different malonate ligands at the apices (Table 1 and Fig. 1). In the crystal structure, adjacent Na polyhedra are linked by a shared edge on opposite sides in the equatorial plane to form an infinite one-dimensional chain. These chains are connected by malonate bridges to form two-dimensional layers (Fig. 2). The VO<sub>5</sub>N<sub>2</sub> polyhedra are grafted on to this layer *via* carboxylate atoms O1 and O2 of the malonate ligands, which are distributed on both sides of the layers owing to the

alternating orientation of the two carboxylate groups (Fig. 2). This arrangement model for VO<sub>5</sub>N<sub>2</sub> favours the minimization of steric hindrance and boosts the stability of the crystal structure.

The remarkable organization of the crystal structure of (I) can be recognized in a view along the *c* axis (Fig. 3), which shows the two-dimensional layers parallel to the (010) plane. An extensive hydrogen-bonding network (Table 2) links the different layers through different functional groups, such as hydroxylamide O and water O, or hydroxylamide N and carboxyl O. This leads to the formation of a stable three-dimensional supramolecular structure.

Experimental

NH<sub>4</sub>VO<sub>3</sub> (1.375 mmol), malonic acid (2.637 mmol) and NaOH (7.863 mmol) were dissolved in H<sub>2</sub>O (10 ml) at room temperature. The resulting light-yellow solution was stirred for approximately 0.5 h in an ice bath. NH<sub>2</sub>OH·HCl (7.326 mmol) was added gradually with constant stirring for 0.5–1.0 h. The resulting yellow solution (pH = 6.02) was filtered. Colourless crystals of (I) suitable for single-crystal X-ray diffraction were obtained by slow evaporation of a mixture of the filtrate and anhydrous ethanol at 277 K over a period of a few days.

Table 1

Selected geometric parameters (Å, °).

V1—N1	2.0193 (11)	Na1—O1W	2.4305 (13)
V1—N2	2.0080 (12)	Na1—O1W <sup>i</sup>	2.4169 (13)
V1—O1	2.1470 (9)	Na1—O2W	2.4132 (13)
V1—O2	2.0368 (9)	Na1—O2W <sup>ii</sup>	2.4140 (13)
V1—O5	1.9030 (10)	Na1—O3	2.6027 (13)
V1—O6	1.8969 (10)	Na1—O4 <sup>iii</sup>	2.5303 (14)
V1—O7	1.6017 (10)	N1—O5	1.4000 (15)
Na1—O1	2.9173 (12)	N2—O6	1.3969 (15)
O1—V1—O6	84.60 (4)	N1—V1—O5	41.69 (5)
O1—V1—O7	171.06 (5)	N1—V1—O6	128.15 (5)
O2—V1—O5	136.61 (4)	N1—V1—O7	96.99 (5)
O2—V1—O6	132.05 (4)	N2—V1—O2	90.89 (4)
O2—V1—O7	89.58 (5)	N2—V1—O5	128.56 (5)
O5—V1—O6	87.28 (4)	N2—V1—O6	41.80 (5)
O5—V1—O7	100.38 (5)	N2—V1—O7	97.75 (5)
O6—V1—O7	101.96 (5)	O2W—Na1—O3	102.47 (4)
N1—V1—N2	163.98 (5)	O3—Na1—O4 <sup>iii</sup>	145.72 (5)

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

Table 2

Hydrogen-bond geometry (Å, °).

D—H...A	D—H	H...A	D...A	D—H...A
O1W—H1WA...O1	0.85	2.58	3.1245 (15)	123
O1W—H1WB...O5 <sup>iv</sup>	0.85	2.24	2.8987 (14)	134
O1W—H1WA...O6	0.85	2.07	2.8994 (15)	166
O2W—H2WA...O5	0.85	2.03	2.8633 (15)	167
O2W—H2WB...O6 <sup>v</sup>	0.85	2.31	2.9701 (14)	135
N1—H1A...O3 <sup>i</sup>	0.84	2.15	2.9717 (15)	163
N1—H2A...O2 <sup>v</sup>	0.84	2.13	2.9635 (15)	172
N2—H1B...O2 <sup>vi</sup>	0.85	2.10	2.9418 (15)	173
N2—H2B...O3 <sup>ii</sup>	0.85	2.07	2.9010 (15)	168

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

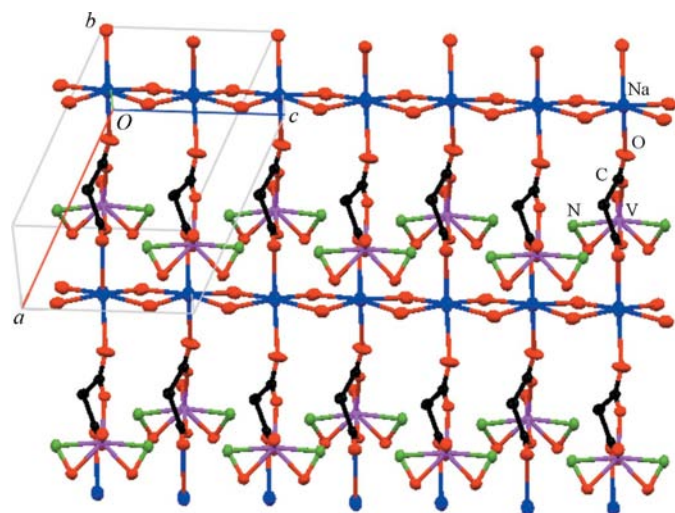


Figure 2

The two-dimensional layer structure of (I), viewed in the *ac* plane. Atom labels represent atoms types and H atoms have been omitted for clarity.

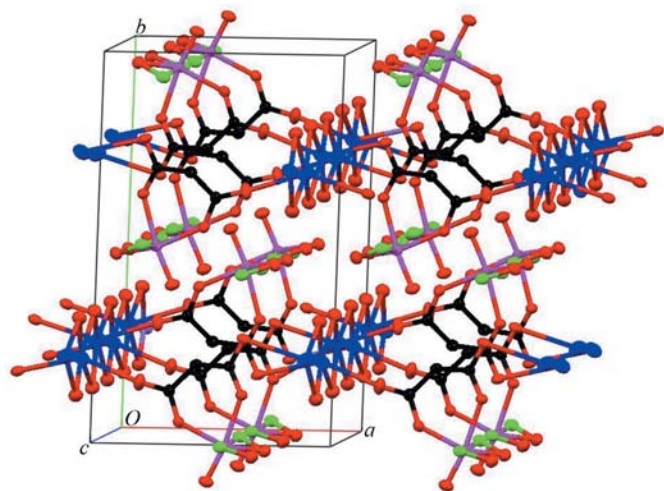


Figure 3

A packing view of (I), showing the two-dimensional layers parallel to the (010) plane. H atoms have been omitted for clarity.

**Table 3**

Selected bond distances (Å) and angles (°) in vanadium(V) hydroxylamide complexes.

Compound	N—O	V—O	V—N	O—V—N	Ref
NH <sub>2</sub> OH	1.47				(a)
[VO(H <sub>2</sub> NO)(C <sub>7</sub> H <sub>3</sub> NO <sub>4</sub> )(H <sub>2</sub> O)]	1.3710 (4)	1.9030 (3)	2.0070 (3)	40.93 (2)	(b)
[VO(H <sub>2</sub> NO) <sub>2</sub> (GlyGly)]·H <sub>2</sub> O	1.397 (3)	1.8961 (14)	2.0046 (16)	41.87 (7)	(c)
[VO(H <sub>2</sub> NO) <sub>2</sub> (GlyGly)]·H <sub>2</sub> O	1.3960 (19)	1.889 (2)	2.0165 (15)	41.72 (7)	
[VO(H <sub>2</sub> NO) <sub>2</sub> (GlyGly)]·H <sub>2</sub> O	1.4040 (3)	1.8920 (3)	2.0210 (4)	41.90 (2)	(d)
[VO(H <sub>2</sub> NO) <sub>2</sub> (Gly)]·H <sub>2</sub> O	1.3970 (4)	1.9080 (3)	2.0070 (4)	41.90 (2)	
[VO(H <sub>2</sub> NO) <sub>2</sub> (Gly)]·H <sub>2</sub> O	1.4050 (3)	1.8980 (4)	2.0180 (3)	41.89 (3)	(d)
[VO(H <sub>2</sub> NO) <sub>2</sub> (Gly)]·H <sub>2</sub> O	1.4020 (3)	1.9020 (3)	2.0100 (4)	41.91 (4)	
[VO(H <sub>2</sub> NO) <sub>2</sub> (Ser)]	1.3980 (5)	1.8990 (5)	2.0100 (3)	41.08 (5)	(d)
[VO(H <sub>2</sub> NO) <sub>2</sub> (Ser)]	1.3870 (4)	1.8940 (4)	2.0040 (3)	41.56 (15)	
[VO(H <sub>2</sub> NO) <sub>2</sub> (imidazole) <sub>2</sub> ]	1.4030 (4)	1.9290 (3)	1.9910 (4)	41.90 (14)	(d)
[VO(H <sub>2</sub> NO) <sub>2</sub> (imidazole) <sub>2</sub> ]	1.3900 (3)	1.9130 (4)	1.9940 (3)	41.63 (9)	
[VO(H <sub>2</sub> NO) <sub>2</sub> (Ala)]·2H <sub>2</sub> O	1.4070 (11)	1.9160 (8)	2.0280 (10)	41.70 (3)	(e)
[VO(H <sub>2</sub> NO) <sub>2</sub> (Ala)]·2H <sub>2</sub> O	1.3830 (10)	1.9080 (9)	1.9970 (10)	41.40 (3)	
[VO(H <sub>2</sub> NO) <sub>2</sub> (Thr)]	1.3980 (3)	1.8960 (2)	2.0140 (3)	41.80 (9)	(e)
[VO(H <sub>2</sub> NO) <sub>2</sub> (Thr)]	1.3940 (4)	1.8830 (2)	2.0270 (3)	41.33 (12)	
Na[VO(NH <sub>2</sub> O) <sub>2</sub> (C <sub>7</sub> H <sub>2</sub> O <sub>4</sub> )]·H <sub>2</sub> O	1.4002 (15)	1.9031 (10)	2.0193 (11)	41.70 (4)	(f)
Na[VO(NH <sub>2</sub> O) <sub>2</sub> (C <sub>7</sub> H <sub>2</sub> O <sub>4</sub> )]·H <sub>2</sub> O	1.3972 (15)	1.8970 (10)	2.0080 (12)	41.81 (5)	

References: (a) Meyers *et al.* (1955); (b) Nuber *et al.* (1981); (c) Paul *et al.* (1997); (d) Keramidis *et al.* (1997); (e) Li *et al.* (2004); (f) this work.**Crystal data**

[NaV(C <sub>3</sub> H <sub>2</sub> O <sub>4</sub> )(NH <sub>2</sub> O) <sub>2</sub> O(H <sub>2</sub> O) <sub>2</sub> ]	$V = 1025.14 (4) \text{ \AA}^3$
$M_r = 292.06$	$Z = 4$
Monoclinic, $P2_1/c$	Mo $K\alpha$ radiation
$a = 9.6393 (2) \text{ \AA}$	$\mu = 1.05 \text{ mm}^{-1}$
$b = 15.4265 (4) \text{ \AA}$	$T = 296 \text{ K}$
$c = 7.4346 (2) \text{ \AA}$	$0.33 \times 0.15 \times 0.13 \text{ mm}$
$\beta = 111.984 (1)^\circ$	

**Data collection**

Bruker APEXII CCD area-detector diffractometer	13757 measured reflections
Absorption correction: multi-scan (SADABS; Sheldrick, 2005)	2529 independent reflections
$T_{\min} = 0.825$ , $T_{\max} = 0.873$	2306 reflections with $I > 2\sigma(I)$
	$R_{\text{int}} = 0.022$

**Refinement**

$R[F^2 > 2\sigma(F^2)] = 0.023$	145 parameters
$wR(F^2) = 0.065$	H-atom parameters constrained
$S = 1.02$	$\Delta\rho_{\text{max}} = 0.36 \text{ e \AA}^{-3}$
2529 reflections	$\Delta\rho_{\text{min}} = -0.30 \text{ e \AA}^{-3}$

All H atoms are placed in calculated positions and refined using a riding model, with  $U_{\text{iso}}(\text{H})$  values of  $1.2U_{\text{eq}}(\text{carrier})$  for NH<sub>2</sub> and CH<sub>2</sub> groups or  $1.5U_{\text{eq}}(\text{carrier})$  for water molecules.

Data collection: APEX2 (Bruker, 2005); cell refinement: APEX2; data reduction: APEX2; program(s) used to solve structure: SHELXS97 (Sheldrick, 2008); program(s) used to refine structure: SHELXL97 (Sheldrick, 2008); molecular graphics: SHELXTL (Sheldrick, 2008); software used to prepare material for publication: SHELXTL.

The authors acknowledge financial support from the Key Project of the Shanghai Science and Technology Committee (grant Nos. 05JC14070, 06DZ05025 and 08JC1408600).

Supplementary data for this paper are available from the IUCr electronic archives (Reference: DN3120). Services for accessing these data are described at the back of the journal.

**References**

- Bruker (2005). APEX2. Bruker AXS Inc., Madison, Wisconsin, USA.
- Keramidas, A. D., Miller, S. M., Anderson, O. P. & Crans, D. C. (1997). *J. Am. Chem. Soc.* **119**, 8901–8915.
- Li, L. Z., Xu, T. & Wang, D. Q. (2004). *J. Chem. Crystallogr.* **34**, 585–590.
- Meyers, E. A. & Lipscomb, W. N. (1955). *Acta Cryst.* **8**, 583–587.
- Nuber, B. & Weiss, J. (1981). *Acta Cryst.* **B37**, 947–948.
- Paul, P. C., Angus-Dunne, S. J., Batchelor, R. J., Einstein, F. W. B. & Tracey, A. S. (1997). *Can. J. Chem.* **75**, 183–191.
- Sheldrick, G. M. (2005). SADABS. Version 2.10. University of Göttingen, Germany.
- Sheldrick, G. M. (2008). *Acta Cryst.* **A64**, 112–122.
- Thompson, K. H., Tsukada, Y., Xu, Z., Battel, M., McNeill, J. H. & Orvig, C. (2002). *Biol. Trace Elem. Res.* **86**, 31–44.
- Tracey, A. S. (2000). *J. Inorg. Biochem.* **80**, 11–16.