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#### Key indicators

Single-crystal X-ray study T = 120 KMean  $\sigma(\text{C-C}) = 0.004 \text{ Å}$  R factor = 0.043 wR factor = 0.105Data-to-parameter ratio = 18.3

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

# (2,2'-Bipyridyl)bis(pentane-2,4-dionato)vanadium(III) perchlorate dichloromethane solvate

The title compound is a stoichiometrically solvated salt,  $[V(C_5H_7O_2)_2(C_{10}H_8N_2)]ClO_4 \cdot CH_2Cl_2$ . The ionic components are linked by three  $C-H \cdot \cdot \cdot O$  hydrogen bonds into chains from which the solvent dichloromethane molecules are pendant, and pairs of antiparallel (inversion-related) chains are linked by a single  $\pi-\pi$  stacking interaction.

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### Comment

The nitrogen heterocycles 1,10-phenanthroline (phen) and 2,2'-bipyridine (bipy) are among the most widely utilized chelating ligands in coordination chemistry (Lever, 2003). We have recently prepared mixed-ligand vanadium(III) complexes containing both pentane-2,4-dionate (also called acetylacetonate; acac) and 1,10-phenanthroline ligands (Kavitha et al., 2006) in order to assess their antidiabetic activity. Although phen and bipy have similar structures, there is a difference in their chelating ability, which has been attributed to the difference in the geometry of the free molecules (Reyzer & Brodbelt, 1999; Oresmaa et al., 2002). The title complex, (I), which contains a 2,2'-bipyridine ligand, has been prepared in order to compare its structure with that of the 1,10-phenanthroline analogue and with the longer term aim of testing its antidiabetic activity.



In the cation, which has approximate twofold rotational symmetry, the V atom is octahedrally coordinated (Table 1) by three bidentate ligands (two acac and one bipy): each cation is thus chiral. The cation in the arbitrarily chosen asymmetric unit (Fig. 1) has a  $\Delta$  configuration, but space-group symmetry generates a racemic mixture of  $\Lambda$  and  $\Delta$  enantiomers. The component species are linked by three independent two-centre C-H···O hydrogen bonds and one planar three-centre C-H···O hydrogen bond (Table 2). One further C-H···O hydrogen bond links the ionic aggregates into a

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### Figure 1

The asymmetric unit of (I), showing 30% displacement ellipsoids (arbitrary spheres for the H atoms). The C-H···O interactions are indicated by dashed lines.



### Figure 2

A stereoview of part of the crystal structure of (I), showing the formation of a  $\pi$ -stacked pair of hydrogen-bonded (dashed lines) chains. For clarity, the dichloromethane molecules have been omitted, as have the H atoms not involved in the hydrogen-bonding motifs shown.

 $C(10)C(11)[R_2^2(7)]$  (Bernstein *et al.*, 1995) chain of rings running parallel to the [010] direction (Fig. 2); the dichloromethane molecules are pendant from this chain.

A single  $\pi$ - $\pi$  stacking interaction links antiparallel pairs of these chains (Fig. 2). The rings (N21/C22-C26) in the cations at (x, y, z) and (1 - x, 1 - y, 1 - z) are strictly parallel, with an interplanar spacing of 3.426 (2) Å: the corresponding ringcentroid separation is 3.751 (2) Å, and the ring offset is 1.527 (2) Å. Similar C-H···O and  $\pi$ - $\pi$  interactions were identified in the structure of the analogous phen complex  $[V(acac)_2(phen)]ClO_4$  (Kavitha *et al.*, 2006). Otherwise, the bond lengths and angles of (I) present no unusual features, and they are very similar to those in the analogous phen complex.

## **Experimental**

A solution of tris(pentane-2,4-dionato)vanadium(III) (0.30 g) and 2,2'-bipyridinium perchlorate (0.22 g) in methanol (30 ml) was heated under reflux for 3 h under an atmosphere of dinitrogen. The mixture was cooled to yield an orange solid which was crystallized by vapour diffusion of light petroleum into a solution in dichloromethane (m.p. 480 K).

Crystal data

 $[V(C_5H_7O_2)_2(C_{10}H_8N_2)]$ - $D_r = 1.529 \text{ Mg m}^{-3}$ ClO<sub>4</sub>·CH<sub>2</sub>Cl<sub>2</sub> Mo  $K\alpha$  radiation  $M_r = 589.71$ Cell parameters from 5869 Monoclinic,  $P2_1/c$ reflections a = 15.0676 (5) Å  $\theta = 3.2 - 27.5^{\circ}$  $\mu = 0.75 \text{ mm}^{-1}$ b = 12.5534 (3) Å c = 14.5533 (5) Å T = 120 (2) K  $\beta = 111.4860 \ (13)^{\circ}$ Plate, orange  $V = 2561.45 (14) \text{ Å}^3$  $0.50 \times 0.20 \times 0.06$  mm Z = 4

### Data collection

Bruker–Nonius KappaCCD	5869 independent reflections
diffractometer	4242 reflections with $I > 2\sigma(I)$
$\varphi$ and $\omega$ scans	$R_{\rm int} = 0.058$
Absorption correction: multi-scan	$\theta_{\rm max} = 27.5^{\circ}$
(SADABS; Sheldrick, 2003)	$h = -19 \rightarrow 19$
$T_{\min} = 0.706, \ T_{\max} = 0.956$	$k = -16 \rightarrow 16$
37694 measured reflections	$l = -18 \rightarrow 18$

### Refinement

Refinement on $F^2$	$w = 1/[\sigma^2(F_o^2) + (0.0354P)^2]$
$R[F^2 > 2\sigma(F^2)] = 0.043$	+ 2.8417 <i>P</i> ]
$wR(F^2) = 0.105$	where $P = (F_0^2 + 2F_c^2)/3$
S = 1.07	$(\Delta/\sigma)_{\rm max} = 0.001$
5869 reflections	$\Delta \rho_{\rm max} = 0.36 \text{ e } \text{\AA}^{-3}$
320 parameters	$\Delta \rho_{\rm min} = -0.48 \text{ e } \text{\AA}^{-3}$
H-atom parameters constrained	

### Table 1

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Selected bond lengths (Å).
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V1-O33	1.9487 (17)	V1-O31	1.9779 (17)
V1-O43	1.9526 (17)	V1-N11	2.116 (2)
V1-O41	1.9694 (16)	V1-N21	2.125 (2)

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

$D - H \cdot \cdot \cdot A$	D-H	$H \cdots A$	$D \cdots A$	$D - H \cdots A$
C16-H16····O4 <sup>i</sup>	0.95	2.51	3.349 (4)	147
C25-H25···O1	0.95	2.41	3.285 (4)	153
C26-H26···O2	0.95	2.47	3.249 (3)	139
C51-H51A····O43	0.99	2.53	3.429 (4)	150
C51−H51B···O2	0.99	2.59	3.404 (4)	140
C51−H51 <i>B</i> ···O3	0.99	2.44	3.397 (4)	161

Symmetry code: (i) x, y + 1, z.

All H atoms were located in a difference map and then treated as riding atoms, with C-H = 0.95 (ring H), 0.98 (methyl H) or 0.99 Å (CH<sub>2</sub>), and with  $U_{iso}(H) = 1.2U_{eq}(C)$  or  $1.5U_{eq}(methyl C)$ .

Data collection: COLLECT (Hooft, 1999); cell refinement: DENZO (Otwinowski & Minor, 1997) and COLLECT; data reduction: *DENZO* and *COLLECT*; program(s) used to solve structure: *OSCAIL* (McArdle, 2003) and *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *OSCAIL* and *SHELXL97* (Sheldrick, 1997); molecular graphics: *PLATON* (Spek, 2003); software used to prepare material for publication: *SHELXL97* and *PRPKAPPA* (Ferguson, 1999).

X-ray data were collected at the EPSRC X-ray Crystallographic Service, University of Southampton, England. The authors thank the staff for all their help and advice.

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