

{2-[(2-Aminocyclohexyl)iminomethyl]-phenolato}dioxidovanadium(V)

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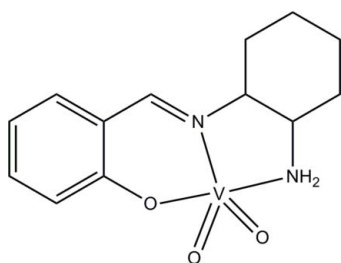
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 Key indicators: single-crystal X-ray study; $T = 298$ K; mean $\sigma(\text{C}-\text{C}) = 0.015$ Å; R factor = 0.099; wR factor = 0.203; data-to-parameter ratio = 14.8.

In the title dioxidovanadium complex, $[\text{V}(\text{C}_{13}\text{H}_{17}\text{N}_2\text{O})\text{O}_2]$, the V^{V} atom is in a square-based pyramidal coordination: the basal plane is defined by the phenolate O, imine N and amine N atoms of the tridentate Schiff base ligand, and by one oxide O atom. The apical position is occupied by the other oxide O atom. In the crystal, molecules are connected by $\text{N}-\text{H}\cdots\text{O}$ and $\text{N}-\text{H}\cdots(\text{O},\text{O})$ hydrogen bonds, forming a tetramer.

Related literature

For related structures and their properties, see: Agarwal & Prasad (2006); Chohan & Sumrra (2010); Huo *et al.* (2004); Jing *et al.* (2005); Lodyga-Chruscinska *et al.* (2008); Xie *et al.* (2004); Yuan *et al.* (2009).



Experimental

Crystal data

$[\text{V}(\text{C}_{13}\text{H}_{17}\text{N}_2\text{O})\text{O}_2]$
 $M_r = 300.23$
 Tetragonal, $I4_1/a$
 $a = 19.120$ (9) Å
 $c = 15.421$ (3) Å
 $V = 5638$ (4) Å³

$Z = 16$
 Mo $K\alpha$ radiation
 $\mu = 0.71$ mm⁻¹
 $T = 298$ K
 $0.23 \times 0.20 \times 0.20$ mm

Data collection

Bruker SMART 1000 CCD diffractometer
 Absorption correction: multi-scan (SADABS; Sheldrick, 2000)
 $T_{\text{min}} = 0.854$, $T_{\text{max}} = 0.871$

19404 measured reflections
 2537 independent reflections
 1487 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.172$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.099$
 $wR(F^2) = 0.203$
 $S = 1.08$
 2537 reflections

172 parameters
 H-atom parameters constrained
 $\Delta\rho_{\text{max}} = 0.44$ e Å⁻³
 $\Delta\rho_{\text{min}} = -0.30$ e Å⁻³

Table 1

Selected bond lengths (Å).

| | | | |
|-------|-----------|-------|-----------|
| V1—O2 | 1.607 (5) | V1—N2 | 2.095 (5) |
| V1—O3 | 1.621 (5) | V1—N1 | 2.137 (6) |
| V1—O1 | 1.900 (5) | | |

Table 2

Hydrogen-bond geometry (Å, °).

| $D-H\cdots A$ | $D-H$ | $H\cdots A$ | $D\cdots A$ | $D-H\cdots A$ |
|---|-------|-------------|-------------|---------------|
| $\text{N2}-\text{H2B}\cdots\text{O2}^{\text{i}}$ | 0.90 | 2.26 | 3.099 (7) | 155 |
| $\text{N2}-\text{H2B}\cdots\text{O3}^{\text{i}}$ | 0.90 | 2.55 | 3.229 (7) | 133 |
| $\text{N2}-\text{H2A}\cdots\text{O3}^{\text{ii}}$ | 0.90 | 2.07 | 2.948 (7) | 166 |

Symmetry codes: (i) $-y + \frac{1}{4}, x + \frac{1}{4}, -z + \frac{5}{4}$; (ii) $y - \frac{1}{4}, -x + \frac{1}{4}, -z + \frac{5}{4}$

Data collection: *SMART* (Bruker, 2000); cell refinement: *SAINT* (Bruker, 2000); data reduction: *SAINT*; program(s) used to solve structure: *SHELXTL* (Sheldrick, 2008); program(s) used to refine structure: *SHELXTL*; molecular graphics: *SHELXTL*; software used to prepare material for publication: *SHELXTL*.

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: HB6676).

References

- Agarwal, R. K. & Prasad, S. (2006). *Rev. Inorg. Chem.* **26**, 471–492.
 Bruker (2000). *SMART* and *SAINT*. Bruker AXS Inc., Madison, Wisconsin, USA.
 Chohan, Z. H. & Sumrra, S. H. (2010). *J. Enzyme Inhib. Med. Chem.* **25**, 599–607.
 Huo, L.-H., Gao, S., Liu, J.-W., Zhao, H. & Ng, S. W. (2004). *Acta Cryst.* **E60**, m606–m608.
 Jing, B., Li, L., Wang, D., Xu, T. & Guo, X. (2005). *Acta Cryst.* **E61**, m2244–m2246.
 Lodyga-Chruscinska, E., Sanna, D., Garribba, E. & Micera, G. (2008). *Dalton Trans.* pp. 4903–4916.
 Sheldrick, G. M. (2000). *SADABS*. University of Göttingen, Germany.
 Sheldrick, G. M. (2008). *Acta Cryst.* **A64**, 112–122.
 Xie, M.-J., Ping, Y.-S., Zheng, L.-D., Hui, J.-Z. & Peng, C. (2004). *Acta Cryst.* **E60**, m1382–m1383.
 Yuan, C. X., Lu, L. P., Gao, X. L., Wu, Y. B., Guo, M. L., Li, Y., Fu, X. Q. & Zhu, M. L. (2009). *J. Biol. Inorg. Chem.* **14**, 841–851.

supplementary materials

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{2-[(2-Aminocyclohexyl)iminomethyl]phenolato}dioxovanadium(V)**Xin-Zhi Sun****Comment**

Oxovanadium complexes with Schiff bases have been received much attention in bioinorganic chemistry (Chohan & Sumrra, 2010; Agarwal & Prasad, 2006; Lodyga-Chruscinska *et al.*, 2008; Yuan *et al.*, 2009). In this paper, the title new dioxovanadium(V) complex, (I), is reported.

In the title dioxovanadium complex, Fig. 1, the V atom is in a square pyramidal coordination. The basal plane of the square pyramid is defined by one phenolate O, one imine N, and one amine N atoms of a Schiff base ligand, and by one oxo O atom. The apical position of the square pyramid is occupied by the other oxo O atom. The bond lengths (Table 1) are comparable to those observed in similar oxovanadium complexes (Xie *et al.*, 2004; Jing *et al.*, 2005; Huo *et al.*, 2004). In the crystal, adjacent molecules are linked through N—H \cdots O and N—H \cdots (O,O) hydrogen bonds (Table 2) to form a tetramer (Fig. 2).

Experimental

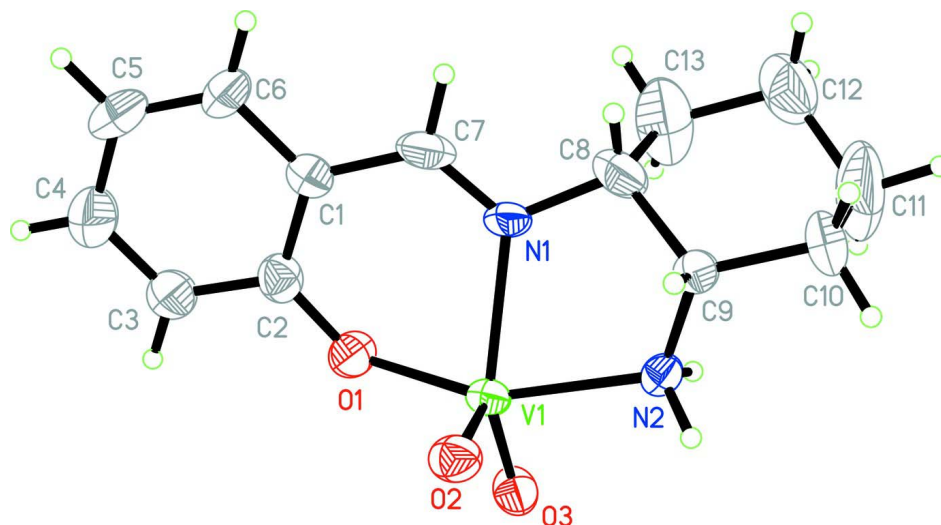
To a MeOH solution (30 ml) of salicylaldehyde (0.122 g, 1.0 mmol) was added a MeOH solution (20 ml) of cyclohexane-1,2-diamine (0.114 g, 1.0 mmol) with stirring. To the above mixture was added a MeOH solution (10 ml) of VO(acac)₂ (0.265 g, 1.0 mmol) with stirring. The mixture was refluxed for 1 h, affording a clear yellow solution. This was allowed to stand at room temperature for a week and block-shaped single crystals were obtained by slow evaporation.

Refinement

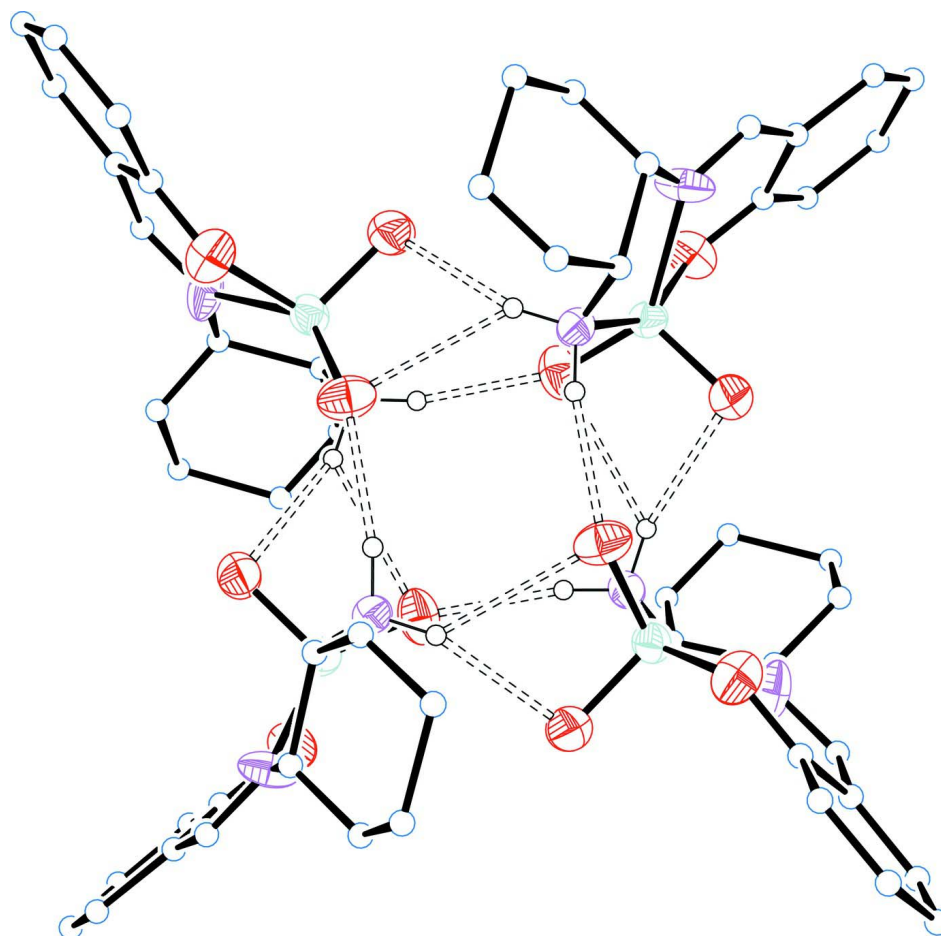
H atoms attached to C and N atoms were placed in geometrically idealized positions with $Csp^2-H = 0.93 \text{ \AA}$, $Csp^3-H = 0.97-0.98 \text{ \AA}$, $N-H = 0.90 \text{ \AA}$, and constrained to ride on their parent atoms, with $U_{iso}(H) = 1.2U_{eq}(C \text{ and } N)$. Apparent disorder was evident in the cyclohexyl ring.

Computing details

Data collection: *SMART* (Bruker, 2000); cell refinement: *SAINTE* (Bruker, 2000); data reduction: *SAINTE* (Bruker, 2000); program(s) used to solve structure: *SHELXTL* (Sheldrick, 2008); program(s) used to refine structure: *SHELXTL* (Sheldrick, 2008); molecular graphics: *SHELXTL* (Sheldrick, 2008); software used to prepare material for publication: *SHELXTL* (Sheldrick, 2008).

**Figure 1**

The structure of the title compound, with displacement ellipsoids drawn at the 30% probability level for Non-H atoms.

**Figure 2**

The tetrameric unit in the crystal, with hydrogen bonds shown as double-dashed lines.

{2-[(2-Aminocyclohexyl)iminomethyl]phenolato}dioxidovanadium(V)

Crystal data

[V(C₁₃H₁₇N₂O)₂]

M_r = 300.23

Tetragonal, *I*₄/a

a = 19.120 (9) Å

c = 15.421 (3) Å

V = 5638 (4) Å³

Z = 16

F(000) = 2496

D_x = 1.415 Mg m⁻³

Mo *Kα* radiation, λ = 0.71073 Å

Cell parameters from 1125 reflections

θ = 2.7–24.5°

μ = 0.71 mm⁻¹

T = 298 K

Block, yellow

0.23 × 0.20 × 0.20 mm

Data collection

Bruker SMART 1000 CCD

diffractometer

Radiation source: fine-focus sealed tube

Graphite monochromator

ω scan

Absorption correction: multi-scan

(*SADABS*; Sheldrick, 2000)

T_{min} = 0.854, *T_{max}* = 0.871

19404 measured reflections

2537 independent reflections

1487 reflections with *I* > 2σ(*I*)

R_{int} = 0.172

θ_{max} = 25.5°, θ_{min} = 1.7°

h = -23→23

k = -23→23

l = -18→18

Refinement

Refinement on *F*²

Least-squares matrix: full

R[*F*² > 2σ(*F*²)] = 0.099

wR(*F*²) = 0.203

S = 1.08

2537 reflections

172 parameters

0 restraints

Primary atom site location: structure-invariant

direct methods

Secondary atom site location: difference Fourier map

Hydrogen site location: inferred from neighbouring sites

H-atom parameters constrained

w = 1/[σ²(*F_o*²) + (0.0708*P*)² + 16.0234*P*]

where *P* = (*F_o*² + 2*F_c*²)/3

(Δ/σ)_{max} < 0.001

Δρ_{max} = 0.44 e Å⁻³

Δρ_{min} = -0.30 e Å⁻³

Special details

Geometry. All e.s.d.'s (except the e.s.d. in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell e.s.d.'s are taken into account individually in the estimation of e.s.d.'s in distances, angles and torsion angles; correlations between e.s.d.'s in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell e.s.d.'s is used for estimating e.s.d.'s involving l.s. planes.

Refinement. Refinement of *F*² against ALL reflections. The weighted *R*-factor *wR* and goodness of fit *S* are based on *F*², conventional *R*-factors *R* are based on *F*, with *F* set to zero for negative *F*². The threshold expression of *F*² > σ(*F*²) is used only for calculating *R*-factors(gt) etc. and is not relevant to the choice of reflections for refinement. *R*-factors based on *F*² are statistically about twice as large as those based on *F*, and *R*-factors based on ALL data will be even larger.

Fractional atomic coordinates and isotropic or equivalent isotropic displacement parameters (Å²)

| | <i>x</i> | <i>y</i> | <i>z</i> | <i>U_{iso}</i> */ <i>U_{eq}</i> |
|-----|-------------|-------------|-------------|---|
| V1 | 0.04762 (6) | 0.08630 (6) | 0.66104 (7) | 0.0417 (4) |
| N1 | 0.1074 (3) | 0.0188 (3) | 0.5792 (4) | 0.0574 (17) |
| N2 | 0.0524 (3) | 0.1379 (3) | 0.5413 (3) | 0.0424 (14) |
| H2A | 0.0122 | 0.1616 | 0.5337 | 0.051* |
| H2B | 0.0872 | 0.1695 | 0.5436 | 0.051* |
| O1 | 0.0854 (3) | 0.0299 (3) | 0.7510 (3) | 0.0589 (14) |

| | | | | |
|------|-------------|-------------|-------------|-------------|
| O2 | -0.0330 (2) | 0.0631 (2) | 0.6543 (3) | 0.0581 (14) |
| O3 | 0.0526 (3) | 0.1602 (2) | 0.7122 (3) | 0.0596 (14) |
| C1 | 0.1142 (4) | -0.0770 (4) | 0.6803 (5) | 0.062 (2) |
| C2 | 0.0962 (4) | -0.0379 (4) | 0.7536 (5) | 0.057 (2) |
| C3 | 0.0907 (5) | -0.0727 (5) | 0.8315 (6) | 0.078 (3) |
| H3 | 0.0788 | -0.0477 | 0.8811 | 0.093* |
| C4 | 0.1022 (6) | -0.1422 (6) | 0.8374 (7) | 0.109 (4) |
| H4 | 0.0984 | -0.1643 | 0.8909 | 0.130* |
| C5 | 0.1195 (6) | -0.1808 (5) | 0.7646 (8) | 0.111 (4) |
| H5 | 0.1271 | -0.2287 | 0.7692 | 0.133* |
| C6 | 0.1251 (5) | -0.1488 (5) | 0.6875 (7) | 0.094 (3) |
| H6 | 0.1365 | -0.1748 | 0.6384 | 0.112* |
| C7 | 0.1238 (4) | -0.0435 (5) | 0.5984 (5) | 0.073 (3) |
| H7 | 0.1441 | -0.0701 | 0.5546 | 0.087* |
| C8 | 0.1201 (5) | 0.0398 (6) | 0.4903 (6) | 0.094 (3) |
| H8 | 0.1107 | -0.0015 | 0.4546 | 0.112* |
| C9 | 0.0635 (4) | 0.0936 (4) | 0.4666 (5) | 0.063 (2) |
| H9 | 0.0202 | 0.0670 | 0.4586 | 0.075* |
| C10 | 0.0774 (7) | 0.1306 (6) | 0.3813 (6) | 0.110 (4) |
| H10A | 0.0630 | 0.0998 | 0.3346 | 0.132* |
| H10B | 0.0478 | 0.1717 | 0.3788 | 0.132* |
| C11 | 0.1431 (8) | 0.1505 (8) | 0.3664 (10) | 0.167 (7) |
| H11A | 0.1534 | 0.1903 | 0.4032 | 0.200* |
| H11B | 0.1465 | 0.1661 | 0.3067 | 0.200* |
| C12 | 0.1999 (6) | 0.0933 (7) | 0.3823 (7) | 0.115 (4) |
| H12A | 0.1972 | 0.0579 | 0.3373 | 0.137* |
| H12B | 0.2461 | 0.1141 | 0.3804 | 0.137* |
| C13 | 0.1873 (5) | 0.0597 (7) | 0.4717 (8) | 0.134 (5) |
| H13A | 0.2170 | 0.0187 | 0.4764 | 0.161* |
| H13B | 0.2023 | 0.0926 | 0.5158 | 0.161* |

Atomic displacement parameters (\AA^2)

| | U^{11} | U^{22} | U^{33} | U^{12} | U^{13} | U^{23} |
|-----|------------|------------|------------|------------|------------|-------------|
| V1 | 0.0364 (7) | 0.0422 (8) | 0.0467 (7) | 0.0001 (6) | 0.0060 (6) | -0.0070 (6) |
| N1 | 0.056 (4) | 0.066 (4) | 0.050 (4) | 0.032 (3) | 0.012 (3) | 0.003 (3) |
| N2 | 0.033 (3) | 0.039 (3) | 0.055 (4) | -0.001 (3) | 0.000 (3) | 0.006 (3) |
| O1 | 0.072 (4) | 0.055 (3) | 0.049 (3) | 0.009 (3) | -0.008 (3) | -0.004 (3) |
| O2 | 0.054 (3) | 0.049 (3) | 0.071 (4) | -0.002 (2) | 0.009 (3) | -0.005 (3) |
| O3 | 0.082 (4) | 0.047 (3) | 0.050 (3) | -0.010 (3) | 0.008 (3) | -0.007 (2) |
| C1 | 0.076 (6) | 0.061 (5) | 0.048 (5) | 0.010 (4) | 0.013 (4) | -0.001 (4) |
| C2 | 0.057 (5) | 0.064 (6) | 0.050 (5) | 0.002 (4) | -0.003 (4) | 0.005 (4) |
| C3 | 0.105 (7) | 0.073 (6) | 0.055 (6) | 0.018 (5) | 0.010 (5) | 0.002 (5) |
| C4 | 0.160 (11) | 0.085 (8) | 0.080 (7) | 0.024 (7) | 0.023 (7) | 0.025 (7) |
| C5 | 0.167 (11) | 0.051 (6) | 0.115 (10) | 0.037 (6) | 0.030 (8) | 0.003 (6) |
| C6 | 0.150 (10) | 0.055 (6) | 0.076 (7) | 0.040 (6) | 0.022 (6) | 0.015 (5) |
| C7 | 0.073 (6) | 0.097 (7) | 0.048 (5) | 0.044 (5) | 0.005 (4) | -0.012 (5) |
| C8 | 0.095 (8) | 0.124 (8) | 0.062 (6) | 0.045 (6) | 0.034 (5) | 0.014 (6) |
| C9 | 0.083 (6) | 0.063 (5) | 0.043 (4) | 0.028 (5) | 0.003 (4) | 0.002 (4) |
| C10 | 0.161 (12) | 0.115 (9) | 0.054 (6) | 0.037 (8) | 0.029 (7) | 0.036 (6) |

| | | | | | | |
|-----|------------|------------|------------|-------------|------------|------------|
| C11 | 0.142 (13) | 0.200 (16) | 0.158 (13) | -0.015 (12) | 0.024 (10) | 0.123 (12) |
| C12 | 0.111 (9) | 0.135 (10) | 0.098 (8) | -0.012 (8) | 0.034 (7) | 0.026 (7) |
| C13 | 0.058 (7) | 0.215 (14) | 0.129 (10) | 0.002 (7) | -0.005 (7) | 0.079 (10) |

Geometric parameters (Å, °)

| | | | |
|------------|------------|---------------|------------|
| V1—O2 | 1.607 (5) | C5—H5 | 0.9300 |
| V1—O3 | 1.621 (5) | C6—H6 | 0.9300 |
| V1—O1 | 1.900 (5) | C7—H7 | 0.9300 |
| V1—N2 | 2.095 (5) | C8—C13 | 1.370 (12) |
| V1—N1 | 2.137 (6) | C8—C9 | 1.536 (11) |
| N1—C7 | 1.267 (9) | C8—H8 | 0.9800 |
| N1—C8 | 1.448 (10) | C9—C10 | 1.517 (11) |
| N2—C9 | 1.447 (8) | C9—H9 | 0.9800 |
| N2—H2A | 0.9000 | C10—C11 | 1.332 (15) |
| N2—H2B | 0.9000 | C10—H10A | 0.9700 |
| O1—C2 | 1.314 (8) | C10—H10B | 0.9700 |
| C1—C6 | 1.394 (11) | C11—C12 | 1.561 (17) |
| C1—C2 | 1.398 (10) | C11—H11A | 0.9700 |
| C1—C7 | 1.428 (11) | C11—H11B | 0.9700 |
| C2—C3 | 1.377 (11) | C12—C13 | 1.541 (13) |
| C3—C4 | 1.350 (12) | C12—H12A | 0.9700 |
| C3—H3 | 0.9300 | C12—H12B | 0.9700 |
| C4—C5 | 1.383 (13) | C13—H13A | 0.9700 |
| C4—H4 | 0.9300 | C13—H13B | 0.9700 |
| C5—C6 | 1.342 (13) | | |
| O2—V1—O3 | 109.2 (3) | N1—C7—H7 | 116.7 |
| O2—V1—O1 | 104.8 (2) | C1—C7—H7 | 116.7 |
| O3—V1—O1 | 96.7 (2) | C13—C8—N1 | 115.6 (9) |
| O2—V1—N2 | 96.6 (2) | C13—C8—C9 | 115.2 (9) |
| O3—V1—N2 | 90.9 (2) | N1—C8—C9 | 107.0 (6) |
| O1—V1—N2 | 153.4 (2) | C13—C8—H8 | 106.1 |
| O2—V1—N1 | 108.0 (2) | N1—C8—H8 | 106.1 |
| O3—V1—N1 | 141.5 (3) | C9—C8—H8 | 106.1 |
| O1—V1—N1 | 83.4 (2) | N2—C9—C10 | 116.3 (7) |
| N2—V1—N1 | 75.0 (2) | N2—C9—C8 | 107.8 (6) |
| C7—N1—C8 | 116.1 (7) | C10—C9—C8 | 113.3 (7) |
| C7—N1—V1 | 124.2 (5) | N2—C9—H9 | 106.2 |
| C8—N1—V1 | 118.8 (5) | C10—C9—H9 | 106.2 |
| C9—N2—V1 | 115.6 (4) | C8—C9—H9 | 106.2 |
| C9—N2—H2A | 108.4 | C11—C10—C9 | 116.6 (11) |
| V1—N2—H2A | 108.4 | C11—C10—H10A | 108.1 |
| C9—N2—H2B | 108.4 | C9—C10—H10A | 108.1 |
| V1—N2—H2B | 108.4 | C11—C10—H10B | 108.1 |
| H2A—N2—H2B | 107.4 | C9—C10—H10B | 108.1 |
| C2—O1—V1 | 129.9 (5) | H10A—C10—H10B | 107.3 |
| C6—C1—C2 | 119.9 (8) | C10—C11—C12 | 115.4 (11) |
| C6—C1—C7 | 119.5 (8) | C10—C11—H11A | 108.4 |
| C2—C1—C7 | 120.5 (7) | C12—C11—H11A | 108.4 |

| | | | |
|----------|------------|---------------|------------|
| O1—C2—C3 | 119.4 (7) | C10—C11—H11B | 108.4 |
| O1—C2—C1 | 122.8 (7) | C12—C11—H11B | 108.4 |
| C3—C2—C1 | 117.9 (8) | H11A—C11—H11B | 107.5 |
| C4—C3—C2 | 121.4 (9) | C13—C12—C11 | 108.9 (10) |
| C4—C3—H3 | 119.3 | C13—C12—H12A | 109.9 |
| C2—C3—H3 | 119.3 | C11—C12—H12A | 109.9 |
| C3—C4—C5 | 120.7 (10) | C13—C12—H12B | 109.9 |
| C3—C4—H4 | 119.7 | C11—C12—H12B | 109.9 |
| C5—C4—H4 | 119.7 | H12A—C12—H12B | 108.3 |
| C6—C5—C4 | 119.7 (9) | C8—C13—C12 | 116.7 (9) |
| C6—C5—H5 | 120.1 | C8—C13—H13A | 108.1 |
| C4—C5—H5 | 120.1 | C12—C13—H13A | 108.1 |
| C5—C6—C1 | 120.4 (9) | C8—C13—H13B | 108.1 |
| C5—C6—H6 | 119.8 | C12—C13—H13B | 108.1 |
| C1—C6—H6 | 119.8 | H13A—C13—H13B | 107.3 |
| N1—C7—C1 | 126.6 (7) | | |

Hydrogen-bond geometry (Å, °)

| <i>D—H...A</i> | <i>D—H</i> | <i>H...A</i> | <i>D...A</i> | <i>D—H...A</i> |
|---------------------------|------------|--------------|--------------|----------------|
| N2—H2B...O2 ⁱ | 0.90 | 2.26 | 3.099 (7) | 155 |
| N2—H2B...O3 ⁱ | 0.90 | 2.55 | 3.229 (7) | 133 |
| N2—H2A...O3 ⁱⁱ | 0.90 | 2.07 | 2.948 (7) | 166 |

Symmetry codes: (i) $-y+1/4, x+1/4, -z+5/4$; (ii) $y-1/4, -x+1/4, -z+5/4$.