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Key indicators

Single-crystal X-ray study T = 296 KMean $\sigma(\text{C-C}) = 0.016 \text{ Å}$ R factor = 0.042 wR factor = 0.099 Data-to-parameter ratio = 15.4

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

{6,6'-Dimethoxy-2,2'-[ethane-1,2-diylbis-(nitrilomethylidyne)]diphenolato}(ethanol)dioxouranium(VI)

In the title compound, $[UO_2(C_{18}H_{18}N_2O_4)(C_2H_6O)]$, the U atom has a distorted pentagonal–bipyramidal geometry with a tetradentate Schiff base and ethanol ligands in the equatorial plane and oxo atoms in the axial positions. Complex molecules are connected by $O-H\cdots O$ and $C-H\cdots O$ interactions, forming dimers in the crystal structure.

Comment

Many Schiff base complexes have potential biological interest in an attempt to mimic the structural and functional features of multimetal active sites. Schiff bases have also found useful applications in the synthesis of molecular and supramolecular polymetallic functional systems exhibiting specific optical or magnetic properties (Salmon *et al.*, 2004). Here, we report the synthesis and crystal structure of the title compound, (I).



The U^{VI} atom has a distorted pentagonal-bipyramidal coordination geometry, with the Schiff base and ethanol ligands in the equatorial plane (Fig. 1). The phenoxide atoms O1 and O4 are displaced by 0.066 (5) and 0.143 (5) Å, respectively, from the plane defined by O1/N1/N2/O4/U1/O5.



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Figure 1

The molecular structure of (I), showing the atom numbering scheme and 50% probability displacement ellipsoids.



Figure 2

The crystal structure of (I). H atoms have been omitted unless they are involved in hydrogen bonds (dashed lines).

The oxo atoms O6 and O7 occupy the axial positions. The U atom is displaced by 0.053 (3) Å from the equatorial plane towards atom O7. The U1-O6 and U1-O7 bond distances (Table 1) are typical for U=O double bonds, close to those reported previously [1.778 (4) and 1.794 (5) Å; Masci et al., 2005].

In the crystal structure, molecules related by an inversion centre are connected by $O5-H5A\cdots O1^{i}$, $O5-H5A\cdots O2^{i}$, $C17 - H17B \cdot \cdot \cdot O1^{i}$ and C19-H19A···O6ⁱ interactions (Table 1), forming dimers having two intersecting $R_2^2(10)$ and $R_2^2(16)$ rings (Fig. 2). Dimers are interlinked along the c axis via $C8 - H8B \cdots O7^{ii}$ interactions.

Experimental

A solution of UO₂(CH₃COO)₂·2H₂O (1 mmol, 0.424 g) in hot ethanol (10 ml) was added to a hot ethanol solution (15 ml) of N,N'ethylenebis(3-methoxysalicylideneimine) (1 mmol, 0.328 g). The mixture was refluxed for 4 h, with stirring. The mixture was then cooled to room temperature and a red powder was precipitated. The product was filtered and washed with ethanol. Crystals of (I) suitable for X-ray diffraction were grown from an acetone/ethanol/n-hexane (1:1:1 v/v/v) solution by slow evaporation over a period of 2 d (yield 79%).

Crystal data

$\begin{bmatrix} U(C_{18}H_{18}N_2O_4)O_2(C_2H_6O) \end{bmatrix}$ $M_r = 642.44$ Orthorhombic, <i>Pbca</i> a = 9.2132 (4) Å b = 23.6595 (10) Å c = 19.7075 (10) Å $U_1 = 4000$ Å	Z = 8 $D_x = 1.984 \text{ Mg m}^{-3}$ Mo K\alpha radiation $\mu = 7.60 \text{ mm}^{-1}$ T = 296 K Plate, red		
V = 4295.8 (3) A ² Data collection Stoe IPDS-2 diffractometer ω scans Absorption correction: integration	$0.43 \times 0.23 \times 0.05$ mm 57109 measured reflections 4219 independent reflections 2789 reflections with $I > 2\sigma(I)$		

 $R_{\rm int} = 0.104$

 $\theta_{\rm max} = 26.0^\circ$

Absorption correction: integration (X-RED32; Stoe & Cie, 2002) $T_{\rm min}=0.104,\;T_{\rm max}=0.597$

Refinement

Refinement on F^2	$w = 1/[\sigma^2(F_o^2) + (0.0393P)^2]$
$R[F^2 > 2\sigma(F^2)] = 0.042$	+ 9.9627 <i>P</i>]
$wR(F^2) = 0.099$	where $P = (F_0^2 + 2F_c^2)/3$
S = 1.10	$(\Delta/\sigma)_{\rm max} < 0.001$
4219 reflections	$\Delta \rho_{\rm max} = 1.26 \text{ e } \text{\AA}^{-3}$
274 parameters	$\Delta \rho_{\rm min} = -0.68 \text{ e } \text{\AA}^{-3}$
H atoms treated by a mixture of	
independent and constrained	
refinement	

Table 1 Selected geometric parameters (Å, °).

U1-07	1.752 (7)	U1-O5	2.471 (6)
U1-O6	1.763 (7)	U1-N1	2.540 (7)
U1-O4	2.253 (6)	U1-N2	2.542 (8)
U1-O1	2.265 (6)		
O7-U1-O6	177.3 (3)	O7-U1-O5	96.1 (3)
O7-U1-O4	87.8 (3)	O6-U1-O5	86.5 (3)
O6-U1-O4	91.7 (3)	O4-U1-N1	135.6 (2)
O7-U1-O1	90.3 (3)	O1-U1-N1	70.1 (2)
O6-U1-O1	91.3 (3)	O5-U1-N1	146.0 (2)
O4-U1-O1	154.2 (2)	O4-U1-N2	69.9 (2)

l able 2			
Hydrogen-bond	geometry	(Å,	°).

$D - H \cdots A$	<i>D</i> -H	$H \cdots A$	$D \cdots A$	$D - \mathbf{H} \cdots A$
$\begin{array}{c} 05 - H5A \cdots 01^{i} \\ 05 - H5A \cdots 02^{i} \\ C8 - H8B \cdots 07^{ii} \\ C17 - H17B \cdots 01^{i} \\ C18 - H18C \cdots 07 \\ C19 - H19A \cdots 06^{i} \end{array}$	0.83 (6)	2.53 (15)	3.059 (9)	124 (15)
	0.83 (6)	2.19 (8)	2.952 (11)	154 (16)
	0.97	2.50	3.183 (11)	127
	0.97	2.45	3.163 (19)	130
	0.96	2.57	3.382 (19)	142
	0.96	2.59	3.423 (13)	145

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

The hydroxy H atom was located in a difference map and refined with an O-H distance restraint of 0.83 (6) Å $[U_{iso}(H) = 1.5U_{eq}(O)]$. All other H atoms were positioned geometrically and treated as riding, with C-H = 0.93-0.97 Å and $U_{iso}(H) = 1.2U_{eq}(C)$ or $1.5U_{eq}$ (methyl C). The highest residual electron density peak is located 1.21 Å from atom U1.

Data collection: X-AREA (Stoe & Cie, 2002); cell refinement: X-AREA; data reduction: X-RED32 (Stoe & Cie, 2002); program(s) used to solve structure: SHELXS97 (Sheldrick, 1997); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: ORTEP-3 for Windows (Farrugia, 1997); software used to prepare material for publication: WinGX (Farrugia, 1999).

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