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Key indicators

Single-crystal X-ray study T = 298 KMean σ (C–C) = 0.009 Å R factor = 0.034 wR factor = 0.092 Data-to-parameter ratio = 12.6

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

Diaquabis(1,10-phenanthroline- $\kappa^2 N, N'$)cadmium(II) pentafluorooxoniobate(V)

The crystal structure of the title compound, $[Cd(C_{12}H_8N_2)_2-(H_2O)_2][NbOF_5]$, consists of Cd^{II} complex cations and Nb^V complex anions. The Cd^{II} and Nb^V atoms both have a distorted octahedral geometry. The Nb-F bond *trans* to the Nb=O bond is significantly longer than the other four Nb-F bonds.

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Comment

Out-of-center 'primary' electronic distortions are inherent to oxide fluoride anions of the early d⁰ transition metals. In the $[NbOF_{5}]^{2-}$ anion, the Nb^V atom deviates from the center of the octahedron toward the oxygen ligand, forming a short Nb=O bond and long *trans* Nb-F bond. The *trans*-directing property of the $[NbOF_{5}]^{2-}$ anion can be exploited when designing new materials that exhibit important structure-dependent properties such as piezoelectricity, second-order nonlinear optical activity and ferroelectricity (Heier *et al.*, 1998; Welk *et al.*, 2002). Here the synthesis, crystal structure and characterization of a mixed-metal complex $[Cd(C_{12}H_8N_2)_2(H_2O)_2]^{2+}\cdot[NbOF_5]^{2-}$, (I), which incorporates the $[NbOF_5]^{2-}$ anion, is reported.



The molecular structure of (I) is shown in Fig. 1. The crystal structure of (I) is built up of Cd^{II} complex cations and Nb^V complex anions. The Cd^{II} atom has an octahedral coordination geometry with two 1,10-phenanthroline (phen) and two water molecules. The Nb^V atom assumes a distorted octahedral coordination with five F atoms and one O atom. The Nb–F bond *trans* to the Nb=O bond is significantly longer than the other four Nb–F bonds in the same anion (Table 1). This feature was also observed in previously reported structures (Halasyamani *et al.*, 1996; Norquist *et al.*, 1999; Izumi *et al.*, 2005). The coordinated water molecules of the Cd^{II} complex cation form hydrogen bonds with the [NbOF₅]^{2–} anion. Some phen H atoms also interact with the [NbOF₅]^{2–} anion *via* C– H…F hydrogen bonds (Table 2).

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Figure 1

The molecular structure of (I), with displacement ellipsoids drawn at the 30% probability level. All H atoms have been omitted.



Figure 2

The molecular packing of (I), viewed along the *a* axis. For clarity, all H atoms have been omitted. Dashed lines indicate $O-H\cdots F$ hydrogenbonding interactions.

Experimental

All reagents were of analytical grade from commercial sources and used without further purification. Nb₂O₅ (0.133 g, 0.5 mmol) was first dissolved in HF solution (1 ml, 42 wt% in H₂O) at 383 K for 2 h in a Teflon-lined stainless steel vessel. After the solution had been cooled to room temperature, $3CdSO_4$ ·8H₂O (0.257 g, 0.33 mmol), phen (0.496 g, 2.5 mmol) and water (15 ml) were added. The pH of the mixture was adjusted to about 6 using KOH solution. The mixture was then heated under autogenous hydrothermal conditions at 413 K for 3 d. The mixture was filtered, then the solution was allowed to evaporate slowly. After 5 d, colorless single crystals suitable for X-ray diffraction were obtained. The crystals were dried in air. Elemental analysis: found: C 40.31, H 2.78, N 7.87%; calculated for C₂₄H₂₀CdF₅N₄NbO₃: C 40.34, H 2.82, N 7.85%.

Crystal data

 $\begin{bmatrix} Cd(C_{12}H_8N_2)_2(H_2O)_2 \end{bmatrix} \begin{bmatrix} NbOF_5 \end{bmatrix}$ $M_r = 712.75$ Triclinic, $P\overline{1}$ a = 9.487 (4) Å b = 11.537 (5) Å c = 13.007 (5) Å $\alpha = 99.162$ (6)° $\beta = 109.646$ (6)° $\gamma = 100.661$ (5)° V = 1279.3 (9) Å³

Data collection

Bruker SMART CCD area-detector diffractometer φ and w scans Absorption correction: multi-scan (*SADABS*; Sheldrick, 2002) $T_{min} = 0.650, T_{max} = 0.696$ 6750 measured reflections

Refinement

Refinement on F^2 $R[F^2 > 2\sigma(F^2)] = 0.034$ $wR(F^2) = 0.092$ S = 1.004464 reflections 355 parameters H atoms treated by a mixture of independent and constrained refinement

Mo K α radiation Cell parameters from 2883 reflections $\theta = 2.2-27.2^{\circ}$ $\mu = 1.35 \text{ mm}^{-1}$ T = 298 (2) K Block, colorless $0.35 \times 0.33 \times 0.29 \text{ mm}$

 $D_x = 1.850 \text{ Mg m}^{-3}$

Z = 2

4464 independent reflections 3413 reflections with $I > 2\sigma(I)$ $R_{int} = 0.016$ $\theta_{max} = 25.0^{\circ}$ $h = -11 \rightarrow 11$ $k = -13 \rightarrow 13$ $l = -15 \rightarrow 12$

$$\begin{split} &w = 1/[\sigma^2(F_{\rm o}^2) + (0.0456P)^2 \\ &+ 1.0033P] \\ &where \ P = (F_{\rm o}^2 + 2F_{\rm c}^2)/3 \\ (\Delta/\sigma)_{\rm max} = 0.001 \\ \Delta\rho_{\rm max} = 0.50 \ {\rm e} \ {\rm \AA}^{-3} \\ \Delta\rho_{\rm min} = -0.79 \ {\rm e} \ {\rm \AA}^{-3} \end{split}$$

Table 1 Selected bond lengths (Å).

Nb1-O3	1.764 (3)	Cd1-O2	2.279 (3)
Nb1-F4	1.882 (3)	Cd1-O1	2.296 (3)
Nb1-F3	1.903 (3)	Cd1-N1	2.317 (4)
Nb1-F2	1.935 (3)	Cd1-N3	2.324 (4)
Nb1-F1	1.966 (3)	Cd1-N4	2.364 (4)
Nb1-F5	2.046 (3)	Cd1-N2	2.364 (4)

Table 2

Hydrogen-bond geometry (Å, °).

$D - H \cdot \cdot \cdot A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdots A$
$O1-H25\cdots O3^{i}$	0.85 (4)	1.83 (4)	2.672 (5)	177 (5)
$O1-H26\cdots F1$	0.84 (3)	1.84 (3)	2.679 (5)	172 (6)
$O2-H27\cdots F5^{ii}$	0.85 (3)	1.74 (3)	2.582 (5)	171 (6)
$O2-H28\cdots F2$	0.83 (3)	2.12 (5)	2.851 (5)	146 (5)
O2−H28···F5	0.83 (3)	2.44 (3)	3.130 (5)	142 (5)
C11-H11···F5 ⁱⁱⁱ	0.93	2.52	3.427 (7)	166
$C12-H12\cdots F2^{iv}$	0.93	2.43	3.299 (7)	156
C13−H13···F3 ⁱⁱ	0.93	2.48	3.089 (7)	123
$C15-H15\cdots F3^{v}$	0.93	2.33	3.224 (8)	161

Symmetry codes: (i) -x + 1, -y, -z + 1; (ii) -x + 1, -y + 1, -z + 1; (iii) x + 1, y, z; (iv) -x + 2, -y + 1, -z + 1; (v) x - 1, y, z - 1.

The water H atoms were located in a difference Fourier map and refined with the O-H bond length restrained to 0.85 (2) Å, and assigned fixed isotropic displacement parameters of 0.08 Å². Other H atoms were placed at calculated positions and refined as riding, with C-H = 0.93 Å and $U_{\rm iso}(\rm H) = 1.2U_{eq}(\rm C)$.

Data collection: *SMART* (Bruker, 2002); cell refinement: *SAINT* (Bruker, 2002); data reduction: *SAINT*; program(s) used to solve

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structure: *SHELXS97* (Sheldrick, 1990); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *XP* (Siemens, 1994); software used to prepare material for publication: *SHELXTL* (Siemens, 1995).

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