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Hu et al. • $[Ho_2(C_2H_3O_2)_6(C_{12}H_8N_2)_2]$ **m3213**

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Key indicators

Single-crystal X-ray study T = 298 KMean σ (C–C) = 0.019 Å R factor = 0.058 wR factor = 0.133 Data-to-parameter ratio = 14.0

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

Tetra- μ -acetato- $\kappa^4 O:O';\kappa^3 O,O':O';\kappa^3 O:O,O'$ bis[(acetato- $\kappa^2 O,O'$)(1,10-phenanthroline- $\kappa^2 N,N'$)holmium(III)]

The dimeric title complex, $[Ho_2(CH_3COO)_6(C_{12}H_8N_2)_2]$, lies about a centre of symmetry. Each cation is coordinated by a chelating 1,10-phenanthroline and two chelating acetato ligands, one of which also bridges the two Ho atoms. Two other bridging acetate ligands complete the coordination of the nine-coordinate holmium(III) atoms.

Comment

Dinuclear lanthanide complexes with 1,10-phenanthroline and acetate ligands have received much attention due to their unique properties, such as luminescence and magnetism (Wang *et al.*, 1999). We report here the structure of the dinuclear centrosymmetric title complex, (I), in which the holmium(III) atoms are nine-coordinate (Fig. 1). Panagiotopoulos *et al.* (1995) have reported the structure of a cerium analogue in which the central cerium(III) atoms also adopt this configuration

CH

Experimental

 $Ho(ClO_4)_3$ · $6H_2O$ (0.063 g, 0.11 mmol) in MeOH (20 ml) was added to a boiling solution of 1,10-phenanthroline (0.018 g, 0.1 mmol) and sodium acetate (0.025 g, 0.3 mmol) in MeOH (50 ml). After refluxing for 2 h, the mixture was cooled to room temperature. Colourless crystals suitable for X-ray diffraction analysis were obtained in 35% yield by diffusion of Et₂O into the mixture over one week. **Caution!** Although no problems were encountered in this work, perchlorate compounds are potentially explosive. They should be prepared in small amounts and handled with care.

Crystal data

$[Ho_2(C_2H_3O_2)_6(C_{12}H_8N_2)_2]$	V = 897.5 (7) Å ³
$M_r = 1044.53$	Z = 1
Triclinic, P1	$D_x = 1.933 \text{ Mg m}^{-3}$
a = 8.717 (4) Å	Mo $K\alpha$ radiation
b = 8.926 (4) Å	$\mu = 4.45 \text{ mm}^{-1}$
c = 12.914 (6) Å	T = 298 (2) K
$\alpha = 103.318 \ (7)^{\circ}$	Block, colorless
$\beta = 109.007 \ (6)^{\circ}$	$0.05 \times 0.03 \times 0.03$ mm
$\gamma = 98.380 \ (7)^{\circ}$	

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Data collection

Bruker SMART CCD area-detector diffractometer φ and ω scans Absorption correction: multi-scan (SADABS; Bruker, 2000) $T_{\min} = 0.855, T_{\max} = 0.873$

Refinement

Refinement on F^2 $R[F^2 > 2\sigma(F^2)] = 0.058$ wR(F²) = 0.133 S = 1.033449 reflections 247 parameters

Table 1

Hydrogen-bond geometry (Å, °).

$D - H \cdots A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - H \cdots A$
C1-H1···O5	0.93	2.53	3.042 (14)	115
C10−H10···O6	0.93	2.41	3.031 (14)	124
$C15-H15A\cdots O2^{i}$	0.96	2.46	3.419 (14)	174

6456 measured reflections

 $R_{\rm int}=0.054$

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

3449 independent reflections

2749 reflections with $I > 2\sigma(I)$

H-atom parameters constrained

 $w = 1/[\sigma^2(F_o^2) + (0.0675P)^2]$ where $P = (F_o^2 + 2F_c^2)/3$

 $(\Delta/\sigma)_{\rm max} < 0.001$

 $\Delta \rho_{\rm max} = 2.03 \text{ e} \text{ Å}^{-3}$

 $\Delta \rho_{\rm min} = -1.86 \text{ e } \text{\AA}^{-3}$

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

All H atoms were positioned geometrically and refined using a riding model, with C-H = 0.93 Å and $U_{iso}(H) = 1.2U_{eq}(C)$ for aromatic, and C-H = 0.96 Å and $U_{iso}(H) = 1.5U_{eq}(C)$ for methyl H atoms. The highest residual density peak is located 1.07 Å from the Ho atom and the deepest hole is located 1.71 Å from atom C5.

Data collection: SMART (Bruker, 2000); cell refinement: SAINT (Bruker, 2000); data reduction: SAINT; program(s) used to solve structure: SHELXTL (Bruker, 2000); program(s) used to refine





The molecular structure of (I), with displacement ellipsoids drawn at the 30% probability level. H atoms have been omitted for clarity. [Symmetry code: (i) 2 - x, -y, -z.]

structure: SHELXTL; molecular graphics: SHELXTL; software used to prepare material for publication: SHELXTL.

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