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#### **Key indicators**

Single-crystal X-ray study T = 298 KMean  $\sigma$ (C–C) = 0.014 Å R factor = 0.033 wR factor = 0.079 Data-to-parameter ratio = 11.6

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

# Tris(nitrato- $\kappa^2 O, O'$ )bis(1,10-phenanthroline- $\kappa^2 N, N'$ )ytterbium(III)

In the title compound,  $[Yb(NO_3)_3(C_{12}H_8N_2)_2]$ , two bidentate 1,10-phenanthroline ligands and three bidentate nitrate anions are coordinated to the Yb cation. A crystallographic twofold rotation axis passes through the Yb atom and one nitrate group. The molecules are linked into a three-dimensional network by five  $C-H\cdots O$  hydrogen bonds.

#### Comment

1,10-Phenanthroline (phen) is a bidentate chelating reagent which can not only act as a terminal ligand but also, potentially, give rise to supramolecular interactions such as aromatic stacking to form interesting structures (Marinescu *et al.*, 2005; Chen & Liu, 2002). 1-Hydroxypyridine-2(1H)-thionato-*S*,*O* (pyrithione) and its metal derivatives have found numerous biochemical applications (Lobana *et al.*, 1999) and are widely used as fungicidal materials. As part of our investigation of the reactions between pyrithione and phen with metals, we attempted to design and synthesize a mixed-ligand rare earth metal complex. However, the crystalline product we isolated was the title compound, (I) (Fig. 1), a complex of ytterbium(III) containing no pyrithione ligand.



In complex (I), the Yb<sup>III</sup> centre is ten-coordinated by four N atoms of two chelating phen ligands and six O donors of three nitrate anions. A crystallographic twofold rotation axis passes through Yb, N2 and O5.

In the crystal structure of (I), the molecules are linked into sheets parallel to the (010) plane by  $C-H\cdots O$  hydrogen bonds, generating  $R_2^2(12)$  and  $R_1^2(4)$  rings (Bernstein *et al.*, 1995) (Fig. 2). Neighbouring sheets are connected by two C- $H\cdots O$  hydrogen bonds (Fig. 3), resulting in a three-dimensional network structure.

#### Experimental

© 2007 International Union of Crystallography All rights reserved A solution of equimolar quantities (2 mmol) of sodium pyrithione and 1,10-phenanthroline in ethanol (20 ml) was stirred for 20 min, Received 18 November 2006 Accepted 27 November 2006





The molecular structure of (I), showing the atom-labelling scheme. Displacement ellipsoids are drawn at the 30% probability level. Unlabelled atoms are related to labelled atoms by the symmetry operator -x + 1, y,  $-z + \frac{3}{2}$ 

and then a solution of ytterbium(III) nitrate (1 mmol) in ethanol (10 ml) was added. The reaction mixture was stirred continuously for 2 h at room temperature and then filtered. X-ray quality crystals of (I) were obtained by evaporation of an aqueous solution over a period of about one month.



#### Figure 2

A portion of the crystal struture of (I), showing the formation of a hydrogen-bonded sheet built up from  $C-H\cdots O$  hydrogen bonds (dashed lines). For clarity, H atoms not involved in hydrogen bonding have been omitted. [Symmetry codes: (A)  $1 - x, y, \frac{3}{2} - z$ ; (B)  $2 - x, y, \frac{3}{2} - z$ ; (C)  $-\frac{1}{2} + x, \frac{3}{2} - y, -\frac{1}{2} + z$ ; (D)  $\frac{1}{2} - x, \frac{3}{2} - y, 1 - z$ ; (E)  $\frac{3}{2} - x, \frac{3}{2} - y, 1 - z$ ; (F) 1 - x, y, z; (G)  $\frac{1}{2} + x, \frac{3}{2} - y, -\frac{1}{2} + z$ .]



#### Figure 3

A different view of the crystal structure of (I), showing the formation of a three-dimensional network structure built up from  $C-H \cdots O$  hydrogen bonds (dashed lines). For clarity, H atoms not involved in hydrogen bonding have been omitted. [Symmetry codes: (A) 1 - x, y,  $\frac{3}{2} - z$ ; (C)  $-\frac{1}{2} + x$ ,  $\frac{3}{2} - y$ ,  $-\frac{1}{2} + z$ ; (D)  $\frac{1}{2} - x$ ,  $\frac{3}{2} - y$ , 1 - z; (H)  $\frac{1}{2} + x$ ,  $-\frac{1}{2} + y$ , z; (I)  $\frac{3}{2} - x$ ,  $-\frac{1}{2} + y$ ,  $\frac{3}{2} - z$ ; (J) 1 - x, 1 - y, 1 - z; (K) x, 1 - y,  $-\frac{1}{2} + z$ .]

Z = 4

 $D_x = 1.916 \text{ Mg m}^{-3}$ Mo K\alpha radiation  $\mu = 3.82 \text{ mm}^{-1}$ T = 298 (2) KBlock, yellow  $0.46 \times 0.16 \times 0.10 \text{ mm}$ 

6189 measured reflections

 $R_{\rm int} = 0.040$ 

 $\theta_{\rm max} = 25.0^{\circ}$ 

2162 independent reflections

1885 reflections with  $I > 2\sigma(I)$ 

Crystal data

$Yb(NO_3)_3(C_{12}H_8N_2)_2]$
$A_r = 719.48$
Aonoclinic, $C2/c$
$= 9.456 (2) \text{ Å}_{1}$
p = 15.451 (3)  Å
= 17.104 (3) Å
$B = 93.687 \ (2)^{\circ}$
$V = 2493.8 (8) \text{ Å}^3$

#### Data collection

Bruker SMART CCD area-detector diffractometer  $\varphi$  and  $\omega$  scans Absorption correction: multi-scan (*SADABS*; Sheldrick, 1996)  $T_{\min} = 0.273, T_{\max} = 0.701$ 

#### Refinement

 $\begin{array}{ll} \mbox{Refinement on } F^2 & w = 1/[\sigma^2(F_o^2) + (0.044P)^2 \\ R[F^2 > 2\sigma(F^2)] = 0.033 & w + 0.0077P] \\ wR(F^2) = 0.079 & where \ P = (F_o^2 + 2F_c^2)/3 \\ S = 1.04 & (\Delta/\sigma)_{max} = 0.001 \\ 2162 \ reflections & \Delta\rho_{max} = 1.00 \ e \ \text{\AA}^{-3} \\ 187 \ parameters & \Delta\rho_{min} = -0.68 \ e \ \text{\AA}^{-3} \end{array}$ 

### Table 1

Selected bond lengths (Å).

Yb1-O4	2.400 (5)	Yb1-O1	2.481 (5)
Yb1-O2	2.451 (4)	Yb1-N3	2.533 (5)
Yb1-N4	2.452 (5)		

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

$D - H \cdot \cdot \cdot A$	D-H	$H \cdots A$	$D \cdots A$	$D - \mathbf{H} \cdot \cdot \cdot A$
C12-H12···O3 <sup>i</sup>	0.93	2.67	3.595 (12)	173
$C12-H12\cdots O2^i$	0.93	2.70	3.412 (10)	134
C9−H9···O1 <sup>ii</sup>	0.93	2.43	3.234 (9)	145
C11-H11···O3 <sup>iii</sup>	0.93	2.53	3.253 (10)	135
$C2{-}H2{\cdot}{\cdot}{\cdot}O4^{iv}$	0.93	2.71	3.309 (11)	123

Symmetry codes: (i)  $x + \frac{1}{2}, -y + \frac{3}{2}, z + \frac{1}{2}$ ; (ii)  $-x + 2, y, -z + \frac{3}{2}$ ; (iii)  $-x + \frac{3}{2}, y + \frac{1}{2}, -z + \frac{3}{2}$ ; (iv)  $-x + \frac{1}{2}, y + \frac{1}{2}, -z + \frac{3}{2}$ ;

All H atoms were located in difference Fourier maps and were subsequently treated as riding atoms, with C–H distances of 0.93 Å and with  $U_{iso}(H) = 1.2U_{eq}(C)$ .

Data collection: *SMART* (Siemens, 1996); cell refinement: *SAINT* (Siemens, 1996); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997*a*); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997*a*); molecular graphics:

*SHELXTL* (Sheldrick, 1997*b*); software used to prepare material for publication: *SHELXTL*.

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