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Key indicators

Single-crystal X-ray study T = 295 KMean $\sigma(O-C) = 0.006 \text{ Å}$ Disorder in main residue R factor = 0.039 wR factor = 0.089 Data-to-parameter ratio = 12.0

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

catena-Poly[[diaqua(2-chloronicotinato- $\kappa^2 O, O'$)erbium(III)]-di- μ -2-chloronicotinato- $\kappa^4 O: O'$]

The asymmetric unit of the title compound, $[Er(C_6H_3Cl-NO_2)_3(H_2O)_2]_n$, has two independent disordered 2-chloronicotinate groups, one of which lies on a twofold rotation axis. The group lying on this symmetry element chelates to the water-coordinated Er atom, which also lies on the rotation axis. Two more 2-chloronicotinate groups bridge two diaqua(2-chloronicotinato)erbium species into a linear chain. The Er atom exists in a square-antiprismatic environment made up of O atoms. Received 27 September 2005 Accepted 14 October 2005 Online 19 October 2005

Comment

Lanthanide derivatives of the three isomeric pyridylcarboxylic acids represent a class of metal pyridylcarboxylates for which there are relatively few structurally authenticated examples (Mao *et al.*, 1998). With erbium(III), a trinicotinate has been isolated as a dihydrate (Baker, Cook, Hutchison, Leask *et al.*, 1991; Baker, Cook, Hutchison, Martineau *et al.*, 1991), but its crystal structure has not been reported to date. The erbium(III) derivative of 2-chloronicotinic acid exists as the title diaqua compound, (I) (Fig. 1).



The formula unit of (I) has one 2-chloronicotinate anion that chelates to the metal atom; the other two behave as bridging ligands to give rise to a linear chain motif (Fig. 2). The metal atom, located on a twofold axis, exists in a squareantiprismatic coordination polyhedron by O atoms (Fig. 3).

Hydrogen bonds link adjacent chains into a three-dimensional network structure (Table 2).

Experimental

Erbium trichloride hexahydrate (0.76 g, 2 mmol) and 2-chloronicotinic acid (0.95 g, 6 mmol) were dissolved in a small volume of

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A plot of a portion of the chain structure of (I). Displacement ellipsoids are drawn at the 50% probability level and H atoms are shown as small spheres of arbitrary radii. Symmetry codes are as given in Table 1.



Figure 2

A diagram of the chain structure. H atoms have been omitted.

water. Aqueous sodium hydroxide solution (0.1 M) was added dropwise until the pH of the solution was about 5. The solution was filtered and set aside for a month for the growth of pink prismatic crystals of (I). CHN analysis, calculated for C18H13Cl3N3O8Er: C 32.13, H 1.95, N 6.24%.; found: C 32.16, H 1.92, N 6.21%.

Crystal data

$[Er(C_6H_3CINO_2)_3(H_2O)_2]$
$M_r = 672.92$
Monoclinic, C2/c
a = 16.324 (3) Å
b = 14.130(3)Å
c = 9.614 (2) Å
$\beta = 104.05 \ (3)^{\circ}$
V = 2151.2 (8) Å ³
Z = 4

Data collection

Rigaku R-AXIS RAPID IP	
diffractometer	
v scans	
Absorption correction: multi-scan	
(ABSCOR; Higashi, 1995)	
$T_{\min} = 0.170, \ T_{\max} = 0.672$	
0139 measured reflections	

Refinement

Refinement on F^2	$w = 1/[\sigma^2(F_o^2) + (0.035P)^2]$
$R[F^2 > 2\sigma(F^2)] = 0.039$	+ 5.2256P]
$wR(F^2) = 0.089$	where $P = (F_0^2 + 2F_c^2)/2$
S = 1.14	$(\Delta/\sigma)_{\rm max} = 0.001$
2464 reflections	$\Delta \rho_{\rm max} = 1.60 \text{ e } \text{\AA}^{-3}$
205 parameters	$\Delta \rho_{\rm min} = -1.81 \text{ e } \text{\AA}^{-3}$
H-atom parameters constrained	

 $D_x = 2.078 \text{ Mg m}^{-3}$ Mo $K\alpha$ radiation Cell parameters from 9329 reflections $\theta = 3.1 - 27.5^{\circ}$ $\mu = 4.33 \text{ mm}^{-1}$

T = 295 (2) K Prism, pink

 $R_{\rm int} = 0.066$ $\theta_{\rm max} = 27.5^{\circ}$ $h=-21\rightarrow 21$ $k = -18 \rightarrow 18$ $l = -11 \rightarrow 12$

 $0.10 \times 0.10 \times 0.10$ mm

2464 independent reflections 2317 reflections with $I > 2\sigma(I)$

 $2F_{c}^{2})/3$

Table 1

Selected geometric parameters (Å, °).

Er1-01	2.237 (3)	Er1-O3	2.450 (4)
$Er1-O2^{i}$	2.330 (3)	Er1 - O1W	2.381 (4)
$O1 - Er1 - O1^{ii}$	154.3 (2)	O2 ⁱ -Er1-O3	72.4 (1)
$O1 - Er1 - O2^{i}$	106.5 (1)	O2 ⁱ -Er1-O3 ⁱⁱ	75.3 (2)
$O1 - Er1 - O2^{iii}$	81.6 (1)	$O2^{i}-Er1-O1W$	140.4 (2)
O1-Er1-O3	77.5 (1)	$O2^{i}-Er1-O1W^{ii}$	74.4 (2)
$O1 - Er1 - O3^{ii}$	128.0 (1)	O3-Er1-O3 ⁱⁱ	53.0 (2)
O1 - Er1 - O1W	84.9 (2)	O3-Er1-O1W	146.8 (1)
$O1 - Er1 - O1W^{ii}$	74.3 (1)	$O3-Er1-O1W^{ii}$	127.3 (1)
$O2^{i}-Er1-O2^{iii}$	143.8 (2)	$O1W$ -Er1- $O1W^{ii}$	72.6 (2)
Symmetry codes: (i) $-x + 1, -y + 1,$	-z+1; (ii) $-x+1,$	$y, -z + \frac{3}{2};$ (iii)
$x, -y + 1, z + \frac{1}{2}$			2

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

$D - H \cdot \cdot \cdot A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdot \cdot \cdot A$
$O1W - H1W1 \cdots N1^{iv}$	0.82	2.24	2.87 (3)	134
$O1W - H1W1 \cdots N1^{iv}$	0.82	2.34	2.95 (3)	132
$O1W-H1W2\cdots O3^{iii}$	0.82	2.02	2.776 (5)	153

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

Both 2-chloronicotinate groups, one lying on a general position and the other on the twofold rotation axis, are disordered. For that on a general position, the disorder affects the pyridyl portion but not the carboxylate -CO2 portion. The bond distances between the two portions were restrained to within 0.01 Å of each other; pairs of distances (e.g. C1-Cl1 and C1'-Cl1', C1-C2 and C1'-C2', etc.) were similarly restrained. The pyridyl rings were restrained to be nearly planar, and the displacement parameters of the primed atoms were set to those of the unprimed atoms. The displacement parameters of the C atoms were restrained to be nearly isotropic. The occupancies refined to 53 (1):47 (1) for the unprimed and primed groups, respectively. For the 2-chloronicotinate group lying on the



Figure 3

A diagram illustrating the eight-coordinate geometry of Er. Symmetry codes are as given in Table 1.

twofold rotation axis, the chloropyridyl portion was allowed to refine over this symmetry element with no atoms constrained to lie on the rotation axis. All atoms in this portion had half-site occupancy. H atoms were positioned geometrically, with C-H = 0.93 Å and $U_{\rm iso}({\rm H}) = 1.2U_{\rm eq}({\rm C})$, and were included in the refinement in the riding-model approximation. The two water H atoms were also positioned geometrically, with O-H = 0.82 Å and $U_{\rm iso}({\rm H}) = 1.2U_{\rm eq}({\rm O})$ and refined as riding. The final difference Fourier map had a large peak/hole at about 1 Å from Er1.

Data collection: *RAPID AUTO* (Rigaku, 1998); cell refinement: *RAPID AUTO*; data reduction: *CrystalStructure* (Rigaku/MSC, 2002); program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *ORTEPII* (Johnson, 1976); software used to prepare material for publication: *SHELXL97*.

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