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Bis[μ -*N*-(2-oxidobenzylidene)pyridine-2-carbohydrazidato]bis[chlorido(methanol- κ O)erbium(III)]

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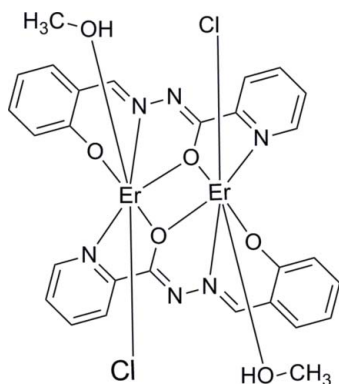
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Key indicators: single-crystal X-ray study; $T = 128$ K; mean $\sigma(\text{C}-\text{C}) = 0.006$ Å; R factor = 0.027; wR factor = 0.059; data-to-parameter ratio = 18.7.

In the binuclear title complex, $[\text{Er}_2(\text{C}_{13}\text{H}_9\text{N}_3\text{O}_2)_2\text{Cl}_2(\text{CH}_3\text{-OH})_2]$, the entire molecule is generated by the application of inversion symmetry. Each Er^{III} ion is seven-coordinated by two O atoms and one N atom from one *N*-(2-oxidobenzylidene)pyridine-2-carbohydrazidate (L^{2-}) ligand, one O atom and one N atom from the symmetry-related L^{2-} ligand, one O atom of a methanol molecule and one chloride anion. The coordination geometry is based on a pseudo-pentagonal bipyramid. Linear supramolecular chains along [010] are formed in the crystal packing through $\text{O}-\text{H}\cdots\text{Cl}$ hydrogen bonds.

Related literature

For complexes containing salicylaldehyde-2-pyridinecarboxylhydrazone and related ligands, see: Guo *et al.* (2011*a,b*); Bai *et al.* (2005, 2006); Wu *et al.* (2004); Milway *et al.* (2003). For the mechanism of the hydrolysis of salicylaldehyde thiosemicarbazone, see: Narang & Aggarwal (1974).



Experimental

Crystal data

 $[\text{Er}_2(\text{C}_{13}\text{H}_9\text{N}_3\text{O}_2)_2\text{Cl}_2(\text{CH}_3\text{O})_2]$ $M_r = 947.97$ Monoclinic, $P2_1/c$ $a = 9.5810$ (4) Å $b = 7.0906$ (3) Å $c = 22.3504$ (8) Å $\beta = 96.920$ (3)° $V = 1507.31$ (10) Å³ $Z = 2$ Mo $K\alpha$ radiation $\mu = 5.76$ mm⁻¹ $T = 128$ K $0.15 \times 0.13 \times 0.12$ mm

Data collection

Bruker SMART CCD

diffractometer

Absorption correction: multi-scan

(SADABS; Bruker 2000)

 $T_{\text{min}} = 0.479$, $T_{\text{max}} = 0.545$

14182 measured reflections

3732 independent reflections

2931 reflections with $I > 2\sigma(I)$ $R_{\text{int}} = 0.040$

Refinement

 $R[F^2 > 2\sigma(F^2)] = 0.027$ $wR(F^2) = 0.059$ $S = 0.99$

3732 reflections

200 parameters

H-atom parameters constrained

 $\Delta\rho_{\text{max}} = 1.25$ e Å⁻³ $\Delta\rho_{\text{min}} = -0.71$ e Å⁻³

Table 1

Selected bond lengths (Å).

Er1—O1	2.157 (3)	Er1—N3	2.433 (3)
Er1—O2 ⁱ	2.284 (3)	Er1—N1	2.488 (3)
Er1—O2	2.316 (3)	Er1—Cl1	2.5901 (12)
Er1—O3	2.327 (3)		

Symmetry code: (i) $-x + 1, -y + 2, -z + 1$.

Table 2

Hydrogen-bond geometry (Å, °).

$D-\text{H}\cdots A$	$D-\text{H}$	$\text{H}\cdots A$	$D\cdots A$	$D-\text{H}\cdots A$
$\text{O3}-\text{H7}\cdots\text{Cl1}^{\text{ii}}$	0.95	2.42	3.128 (4)	131

Symmetry code: (ii) $x, y + 1, z$.

Data collection: SMART (Bruker, 2000); cell refinement: SAINT (Bruker, 2000); data reduction: SAINT; program(s) used to solve structure: SHELXS97 (Sheldrick, 2008); program(s) used to refine structure: SHELXL97 (Sheldrick, 2008); molecular graphics: SHELXTL (Sheldrick, 2008); software used to prepare material for publication: SHELXTL.

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: TK5069).

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supplementary materials

Acta Cryst. (2012). E68, m580–m581 [doi:10.1107/S1600536812013979]

**Bis[μ -*N*-(2-oxidobenzylidene)pyridine-2-carbohydrazidato]bis-
[chlorido(methanol- κ O)erbium(III)]****Hua Yang****Comment**

The chemistry of coordination complexes supported by salicylaldehyde-2-pyridinecarboxyl-hydrazone (H_2L) and its derivatives has received intensive attention as these form coordination complexes with aesthetically pleasing structures and intriguing magnetic behaviour (Guo *et al.*, 2011*a,b*). A handful of transition metal complexes based on the H_2L ligand have been prepared (Bai *et al.*, 2005; Wu *et al.*, 2004; Bai *et al.*, 2006; Milway *et al.*, 2003), but no complex containing rare earth elements has been reported to date. Herein, we report the structure of a new dinuclear Er^{III} complex (Scheme 1). The complex was synthesized by the 2:1:1 reaction of $ErCl_3 \cdot 6H_2O/\alpha$ -pyridoin/salicylaldehyde thiosemicarbazone under solvothermal conditions. The X-ray analysis reveals that the centrosymmetric complex consists of two Er^{III} ions, two L^{2-} ligands, two Cl^- ions and two methanol molecules, Fig. 1 and Table 1. The intermolecular O—H \cdots Cl hydrogen bonds, Table 2, lead to linear supramolecular chains along [010] (Fig. 2).

The remarkable structural feature of the complex is the presence of the *in situ* formed H_2L ligand, which was proposed to be constructed by the reaction of picolinic acid, hydrazine and salicylaldehyde. The picolinic acid was assumed to be derived from the hydrolysis of α -pyridoin, and hydrazine and salicylaldehyde were believed to be originated from the hydrolysis of salicylaldehyde thiosemicarbazone (Narang & Aggarwal, 1974).

Experimental

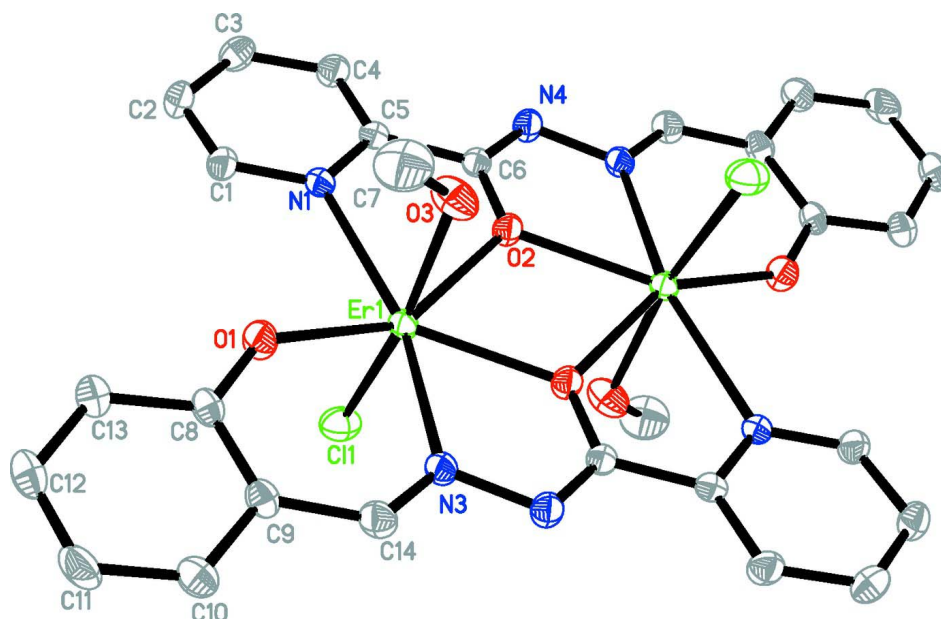
A mixture of $ErCl_3 \cdot 6H_2O$ (0.0762 g, 0.2 mmol), α -pyridoin (0.0214 g, 0.1 mmol), salicylaldehyde thiosemicarbazone (0.0390 g, 0.2 mmol) and CH_3OH (2 ml) was sealed in a 6 ml Pyrex-tube. The tube was heated at 393 K for 3 days under autogenous pressure. Cooling of the resultant solution to room temperature gave yellow crystals. The crystals were collected by filtration, washed with CH_3OH (2 ml) and dried in air.

Refinement

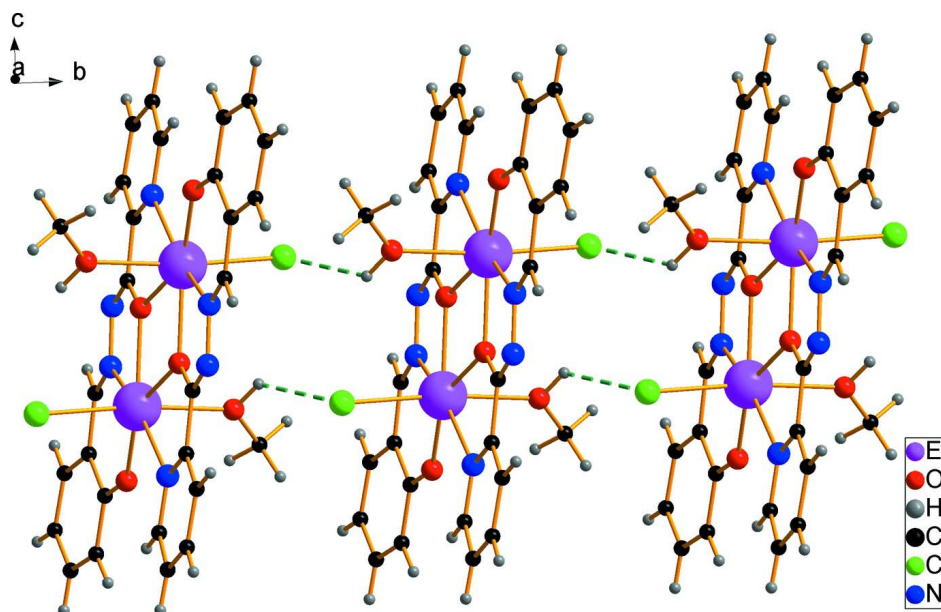
The H atoms were placed in calculated positions with O—H = 0.95 Å and C—H = 0.95–0.98 Å, and with $U_{iso}(H) = 1.2$ – $1.5U_{eq}(C, O)$.

Computing details

Data collection: *SMART* (Bruker, 2000); cell refinement: *SAINTE* (Bruker, 2000); data reduction: *SAINTE* (Bruker, 2000); program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *SHELXTL* (Sheldrick, 2008); software used to prepare material for publication: *SHELXTL* (Sheldrick, 2008).


Figure 1

The molecular structure of the title compound, with atom labels and 30% probability displacement ellipsoids. The H atoms have been omitted for clarity.


Figure 2

View of the linear supramolecular chain along [010] with the O—H...Cl hydrogen bonds shown as dashed lines.

Bis[μ -N-(2-oxido-benzylidene)pyridine-2-carbohydrazidato]bis[chlorido(methanol- κ O)erbium(III)]

Crystal data

[Er₂(C₁₃H₉N₃O₂)₂Cl₂(CH₄O)₂]

$M_r = 947.97$

Monoclinic, $P2_1/c$

Hall symbol: -P 2yc

$a = 9.5810(4) \text{ \AA}$

$b = 7.0906(3) \text{ \AA}$

$c = 22.3504 (8) \text{ \AA}$
 $\beta = 96.920 (3)^\circ$
 $V = 1507.31 (10) \text{ \AA}^3$
 $Z = 2$
 $F(000) = 908$
 $D_x = 2.089 \text{ Mg m}^{-3}$
 Mo $K\alpha$ radiation, $\lambda = 0.71073 \text{ \AA}$

Cell parameters from 4466 reflections

$\theta = 2.7\text{--}28.1^\circ$
 $\mu = 5.76 \text{ mm}^{-1}$
 $T = 128 \text{ K}$
 Block, yellow
 $0.15 \times 0.13 \times 0.12 \text{ mm}$

Data collection

Bruker SMART CCD
 diffractometer
 Radiation source: fine-focus sealed tube
 Graphite monochromator
 φ scans and ω scans
 Absorption correction: multi-scan
 (SADABS; Bruker 2000)
 $T_{\min} = 0.479$, $T_{\max} = 0.545$

14182 measured reflections
 3732 independent reflections
 2931 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.040$
 $\theta_{\max} = 28.3^\circ$, $\theta_{\min} = 2.1^\circ$
 $h = -12 \rightarrow 11$
 $k = -9 \rightarrow 9$
 $l = -29 \rightarrow 29$

Refinement

Refinement on F^2
 Least-squares matrix: full
 $R[F^2 > 2\sigma(F^2)] = 0.027$
 $wR(F^2) = 0.059$
 $S = 0.99$
 3732 reflections
 200 parameters
 0 restraints
 Primary atom site location: structure-invariant
 direct methods

Secondary atom site location: difference Fourier
 map
 Hydrogen site location: inferred from
 neighbouring sites
 H-atom parameters constrained
 $w = 1/[\sigma^2(F_o^2) + (0.0299P)^2]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} = 0.004$
 $\Delta\rho_{\max} = 1.25 \text{ e \AA}^{-3}$
 $\Delta\rho_{\min} = -0.71 \text{ e \AA}^{-3}$

Special details

Geometry. All e.s.d.'s (except the e.s.d. in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell e.s.d.'s are taken into account individually in the estimation of e.s.d.'s in distances, angles and torsion angles; correlations between e.s.d.'s in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell e.s.d.'s is used for estimating e.s.d.'s involving l.s. planes.

Refinement. Refinement of F^2 against ALL reflections. The weighted R -factor wR and goodness of fit S are based on F^2 , conventional R -factors R are based on F , with F set to zero for negative F^2 . The threshold expression of $F^2 > \sigma(F^2)$ is used only for calculating R -factors(gt) etc. and is not relevant to the choice of reflections for refinement. R -factors based on F^2 are statistically about twice as large as those based on F , and R -factors based on ALL data will be even larger.

Fractional atomic coordinates and isotropic or equivalent isotropic displacement parameters (\AA^2)

	x	y	z	$U_{\text{iso}}^*/U_{\text{eq}}$
Er1	0.372121 (17)	0.91591 (3)	0.432040 (7)	0.02740 (6)
O2	0.5836 (3)	1.0615 (4)	0.46548 (11)	0.0323 (6)
O3	0.2988 (3)	1.2290 (5)	0.42893 (15)	0.0526 (8)
H7	0.3591	1.3150	0.4521	0.063*
C5	0.6425 (4)	1.0757 (5)	0.36561 (16)	0.0289 (8)
Cl1	0.48595 (13)	0.58790 (16)	0.42195 (5)	0.0476 (3)
N1	0.5108 (3)	1.0104 (5)	0.34970 (14)	0.0282 (7)
O1	0.1993 (3)	0.8698 (4)	0.36235 (12)	0.0386 (7)
C14	0.0511 (4)	0.7742 (6)	0.46482 (18)	0.0366 (9)
H14	-0.0122	0.7498	0.4935	0.044*

C8	0.0727 (4)	0.7952 (5)	0.35398 (17)	0.0306 (9)
C6	0.6814 (4)	1.1036 (5)	0.43082 (17)	0.0287 (8)
N3	0.1753 (3)	0.8294 (5)	0.48577 (14)	0.0308 (7)
N4	0.8066 (3)	1.1581 (5)	0.45048 (14)	0.0344 (8)
C1	0.4695 (4)	0.9753 (6)	0.29156 (18)	0.0343 (9)
H1	0.3765	0.9309	0.2802	0.041*
C4	0.7340 (4)	1.1077 (6)	0.32293 (18)	0.0373 (10)
H4	0.8257	1.1559	0.3347	0.045*
C10	-0.1348 (5)	0.6619 (7)	0.3905 (2)	0.0435 (11)
H10	-0.1858	0.6313	0.4230	0.052*
C12	-0.1223 (5)	0.6715 (7)	0.2860 (2)	0.0440 (11)
H12	-0.1625	0.6447	0.2459	0.053*
C9	-0.0018 (4)	0.7457 (6)	0.40305 (17)	0.0332 (8)
C3	0.6899 (5)	1.0688 (6)	0.26356 (19)	0.0413 (10)
H3	0.7513	1.0886	0.2338	0.050*
C13	0.0074 (4)	0.7590 (6)	0.29574 (18)	0.0366 (9)
H13	0.0532	0.7954	0.2621	0.044*
C2	0.5561 (5)	1.0008 (7)	0.24736 (18)	0.0393 (10)
H2	0.5241	0.9721	0.2065	0.047*
C11	-0.1934 (5)	0.6230 (7)	0.3332 (2)	0.0499 (12)
H11	-0.2825	0.5629	0.3262	0.060*
C7	0.1756 (6)	1.3080 (8)	0.3970 (3)	0.0722 (17)
H7A	0.1078	1.2075	0.3850	0.108*
H7B	0.2000	1.3728	0.3609	0.108*
H7C	0.1341	1.3985	0.4229	0.108*

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
Er1	0.02276 (9)	0.03156 (11)	0.02796 (10)	-0.00645 (8)	0.00334 (6)	-0.00213 (8)
O2	0.0269 (13)	0.0435 (18)	0.0273 (14)	-0.0087 (12)	0.0062 (10)	-0.0013 (12)
O3	0.0461 (19)	0.0366 (19)	0.073 (2)	0.0010 (15)	-0.0021 (15)	-0.0050 (17)
C5	0.0294 (18)	0.027 (2)	0.0304 (19)	-0.0045 (16)	0.0041 (15)	0.0008 (17)
C11	0.0544 (7)	0.0371 (6)	0.0535 (7)	0.0029 (5)	0.0145 (5)	0.0048 (5)
N1	0.0286 (16)	0.0276 (17)	0.0285 (17)	-0.0043 (13)	0.0038 (13)	-0.0008 (14)
O1	0.0266 (14)	0.054 (2)	0.0347 (16)	-0.0102 (13)	0.0012 (11)	-0.0015 (13)
C14	0.0283 (19)	0.042 (3)	0.040 (2)	-0.0070 (18)	0.0065 (16)	-0.003 (2)
C8	0.0251 (18)	0.032 (2)	0.034 (2)	0.0003 (15)	-0.0017 (15)	-0.0056 (17)
C6	0.0285 (19)	0.031 (2)	0.0273 (19)	-0.0042 (16)	0.0040 (14)	-0.0006 (16)
N3	0.0264 (16)	0.0361 (19)	0.0299 (17)	-0.0075 (14)	0.0033 (13)	-0.0026 (14)
N4	0.0288 (17)	0.046 (2)	0.0289 (18)	-0.0093 (15)	0.0066 (14)	-0.0029 (15)
C1	0.035 (2)	0.038 (3)	0.029 (2)	-0.0038 (17)	0.0004 (16)	0.0010 (18)
C4	0.032 (2)	0.045 (3)	0.035 (2)	-0.0083 (19)	0.0059 (16)	0.003 (2)
C10	0.035 (2)	0.048 (3)	0.046 (3)	-0.012 (2)	0.0013 (19)	-0.002 (2)
C12	0.036 (2)	0.047 (3)	0.046 (3)	-0.003 (2)	-0.0083 (19)	-0.011 (2)
C9	0.0263 (18)	0.036 (2)	0.036 (2)	-0.0036 (18)	0.0003 (15)	-0.0054 (19)
C3	0.041 (2)	0.049 (3)	0.036 (2)	0.002 (2)	0.0135 (18)	0.005 (2)
C13	0.033 (2)	0.040 (2)	0.036 (2)	0.0003 (19)	0.0012 (16)	-0.006 (2)
C2	0.044 (2)	0.047 (3)	0.026 (2)	0.000 (2)	-0.0002 (18)	0.0001 (19)
C11	0.037 (2)	0.050 (3)	0.060 (3)	-0.018 (2)	-0.007 (2)	-0.005 (2)

C7 0.069 (4) 0.058 (4) 0.088 (4) 0.019 (3) 0.003 (3) 0.013 (3)

Geometric parameters (Å, °)

Er1—O1	2.157 (3)	C6—N4	1.286 (5)
Er1—O2 ⁱ	2.284 (3)	N3—N4 ⁱ	1.417 (4)
Er1—O2	2.316 (3)	N4—N3 ⁱ	1.417 (4)
Er1—O3	2.327 (3)	C1—C2	1.376 (6)
Er1—N3	2.433 (3)	C1—H1	0.9500
Er1—N1	2.488 (3)	C4—C3	1.371 (6)
Er1—Cl1	2.5901 (12)	C4—H4	0.9500
O2—C6	1.320 (4)	C10—C11	1.362 (6)
O2—Er1 ⁱ	2.284 (3)	C10—C9	1.403 (5)
O3—C7	1.419 (6)	C10—H10	0.9500
O3—H7	0.9500	C12—C11	1.369 (7)
C5—N1	1.351 (5)	C12—C13	1.382 (6)
C5—C4	1.390 (5)	C12—H12	0.9500
C5—C6	1.474 (5)	C3—C2	1.377 (6)
N1—C1	1.335 (5)	C3—H3	0.9500
O1—C8	1.316 (4)	C13—H13	0.9500
C14—N3	1.286 (5)	C2—H2	0.9500
C14—C9	1.427 (5)	C11—H11	0.9500
C14—H14	0.9500	C7—H7A	0.9800
C8—C13	1.398 (5)	C7—H7B	0.9800
C8—C9	1.423 (5)	C7—H7C	0.9800
O1—Er1—O2 ⁱ	140.21 (10)	O1—C8—C13	120.5 (4)
O1—Er1—O2	149.99 (10)	O1—C8—C9	122.0 (3)
O2 ⁱ —Er1—O2	66.16 (10)	C13—C8—C9	117.5 (3)
O1—Er1—O3	85.42 (12)	N4—C6—O2	124.5 (3)
O2 ⁱ —Er1—O3	88.97 (11)	N4—C6—C5	119.5 (3)
O2—Er1—O3	80.42 (11)	O2—C6—C5	115.9 (3)
O1—Er1—N3	75.24 (10)	C14—N3—N4 ⁱ	112.4 (3)
O2 ⁱ —Er1—N3	65.42 (10)	C14—N3—Er1	129.4 (3)
O2—Er1—N3	130.77 (10)	N4 ⁱ —N3—Er1	118.2 (2)
O3—Er1—N3	90.34 (11)	C6—N4—N3 ⁱ	111.0 (3)
O1—Er1—N1	86.47 (10)	N1—C1—C2	122.7 (4)
O2 ⁱ —Er1—N1	132.20 (10)	N1—C1—H1	118.6
O2—Er1—N1	66.06 (9)	C2—C1—H1	118.6
O3—Er1—N1	84.69 (11)	C3—C4—C5	119.0 (4)
N3—Er1—N1	161.38 (11)	C3—C4—H4	120.5
O1—Er1—Cl1	95.50 (9)	C5—C4—H4	120.5
O2 ⁱ —Er1—Cl1	96.94 (7)	C11—C10—C9	122.4 (4)
O2—Er1—Cl1	93.86 (7)	C11—C10—H10	118.8
O3—Er1—Cl1	169.37 (9)	C9—C10—H10	118.8
N3—Er1—Cl1	100.15 (8)	C11—C12—C13	120.8 (4)
N1—Er1—Cl1	84.80 (8)	C11—C12—H12	119.6
O1—Er1—Er1 ⁱ	167.32 (8)	C13—C12—H12	119.6
O2 ⁱ —Er1—Er1 ⁱ	33.34 (6)	C10—C9—C8	118.5 (4)
O2—Er1—Er1 ⁱ	32.82 (6)	C10—C9—C14	117.5 (4)

O3—Er1—Er1 ⁱ	83.65 (8)	C8—C9—C14	123.9 (3)
N3—Er1—Er1 ⁱ	98.39 (7)	C4—C3—C2	119.6 (4)
N1—Er1—Er1 ⁱ	98.87 (7)	C4—C3—H3	120.2
C11—Er1—Er1 ⁱ	96.43 (3)	C2—C3—H3	120.2
C6—O2—Er1 ⁱ	120.9 (2)	C12—C13—C8	121.5 (4)
C6—O2—Er1	124.5 (2)	C12—C13—H13	119.3
Er1 ⁱ —O2—Er1	113.84 (10)	C8—C13—H13	119.3
C7—O3—Er1	128.4 (3)	C1—C2—C3	118.7 (4)
C7—O3—H7	115.8	C1—C2—H2	120.7
Er1—O3—H7	115.8	C3—C2—H2	120.7
N1—C5—C4	121.5 (4)	C10—C11—C12	119.2 (4)
N1—C5—C6	115.0 (3)	C10—C11—H11	120.4
C4—C5—C6	123.5 (3)	C12—C11—H11	120.4
C1—N1—C5	118.5 (3)	O3—C7—H7A	109.5
C1—N1—Er1	123.2 (3)	O3—C7—H7B	109.5
C5—N1—Er1	117.6 (2)	H7A—C7—H7B	109.5
C8—O1—Er1	141.0 (2)	O3—C7—H7C	109.5
N3—C14—C9	126.9 (4)	H7A—C7—H7C	109.5
N3—C14—H14	116.5	H7B—C7—H7C	109.5
C9—C14—H14	116.5		

Symmetry code: (i) $-x+1, -y+2, -z+1$.

Hydrogen-bond geometry (\AA , $^\circ$)

<i>D</i> —H \cdots <i>A</i>	<i>D</i> —H	H \cdots <i>A</i>	<i>D</i> \cdots <i>A</i>	<i>D</i> —H \cdots <i>A</i>
O3—H7 \cdots C11 ⁱⁱ	0.95	2.42	3.128 (4)	131

Symmetry code: (ii) $x, y+1, z$.