metal-organic papers

Acta Crystallographica Section E Structure Reports Online

ISSN 1600-5368

Zheng-Bo Han* and Yong-Ke He

School of Chemical Science and Engineering, Liaoning University, Shenyang 110036, People's Republic of China

Correspondence e-mail: ceshzb@lnu.edu.cn

Key indicators

Single-crystal X-ray study T = 293 K Mean σ (C–C) = 0.009 Å Disorder in main residue R factor = 0.040 wR factor = 0.090 Data-to-parameter ratio = 11.8

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

catena-Poly[bis[diaquaisonicotinatogadolinium(III)]bis(µ₃-pyridine-3,4-dicarboxylato)]

The title gadolinium(III) coordination polymer, $[Gd(C_6H_4NO_2)(C_7H_3NO_4)(H_2O)_2]_n$, has a pyridine-3,4dicarboxylate-bridged ribbon structure. The isonicotinate anion chelates to the water-coordinated metal atom, which shows ninefold coordination. The isonicotinate unit arises from the decarboxylation of pyridine-3,4-dicarboxylic acid under the hydrothermal conditions of the synthesis.

Comment

In our previous work, two lanthanide coordination polymers, $[Nd(pydc)_2 \cdot H_2O]_n$ and $[Eu(pydc)(ina)(H_2O)_2]_n$ (pydc = pyridine-3,4-dicarboxylate, ina = isonicotinate), were synthesized by treating Nd^{III} or Eu^{III} nitrates with pyridine-3,4-dicarboxylic acid under hydrothermal conditions (Han *et al.*, 2005). Unexpectedly, decarboxylation occurred, and the pydc group was transformed into an ina group in the hydrothermal synthesis of the Eu^{III} coordination polymer. This study has been extended to the other Ln^{III} coordination polymers as the basis for investigating *in situ* decarboxylation. We report here the synthesis and structure of the title gadolinium(III) analogue, (I).



In complex (I), the Gd^{III} ion is coordinated by nine O atoms, five from three pydc ligands, two from one chelating ina group and two from two aqua ligands. The coordination environment around the Gd^{III} ion is a tricapped trigonal prism

© 2006 International Union of Crystallography All rights reserved Received 16 July 2006 Accepted 25 August 2006





The coordination environment of the metal atom in (I). Displacement ellipsoids are drawn at the 30% probability level. H atoms have been omitted. [Symmetry codes: (i) x - 1, y, z; (ii) -x + 2, -y + 1, -z + 1.]

(Fig. 1). Pairs of Gd atoms are bridged by two μ_2 -carboxylate O atoms of the pydc groups to form dinuclear $Gd_2(pydc)_2$ subunits, which are interconnected through the chelating carboxylate groups of the pydc ligands to form an infinite onedimensional ribbon; the $Gd \cdot \cdot \cdot Gd$ distances are 5.954 (3) and 6.433 (3) Å. These chains are further linked via hydrogen bonds (Table 2) between pyridyl N atoms, carboxylate O atoms and aqua ligands to form a three-dimensional supramolecular network.

Experimental

A mixture of Gd(NO₃)₃·6H₂O (0.5 mmol, 0.226 g), pyridine-3,4dicarboxylic acid (0.5 mmol, 0.087 g), NaOH (1 mmol, 0.04 g), and water (10 ml) was placed in a 23 ml Teflon reactor, which was then heated at 453 K for 6 d. The reactor was cooled to room temperature at a rate of 5 K h^{-1} (yield 72%). CHN analysis for C₁₃H₁₁N₂O₈Gd (found/calculated): C 32.65 (32.50), H 2.38 (2.31), N 5.62% (5.83%).

Crystal data

$\begin{bmatrix} Gd(C_6H_4NO_2)(C_7H_3NO_4)(H_2O)_2 \end{bmatrix}$ $M_r = 480.49$ Monoclinic, $P2_1/n$ a = 5.9541 (19) Å b = 25.047 (8) Å c = 11.215 (4) Å $\beta = 100.775$ (7)° V = 1643.0 (9) Å ³	Z = 4 $D_x = 1.943 \text{ Mg m}^{-3}$ Mo K\alpha radiation $\mu = 4.08 \text{ mm}^{-1}$ T = 293 (2) K Block, colourless $0.37 \times 0.23 \times 0.12 \text{ mm}$
Data collection Bruker APEX area-detector	6192 measured reflections

3112 independent reflections 2803 reflections with $I > 2\sigma(I)$

 $R_{\rm int} = 0.031$

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

Bruker APEX area-detector diffractometer φ and ω scans Absorption correction: multi-scan (SADABS; Sheldrick, 1996) $T_{\min} = 0.314, \ T_{\max} = 0.634$

Refinement

Refinement on F^2 $R[F^2 > 2\sigma(F^2)] = 0.040$ $wR(F^2) = 0.090$ S = 1.05 3112 reflections 264 parameters H atoms treated by a mixture of	$w = 1/[\sigma^{2}(F_{o}^{2}) + (0.0338P)^{2} + 7.8772P]$ where $P = (F_{o}^{2} + 2F_{c}^{2})/3$ $(\Delta/\sigma)_{max} < 0.001$ $\Delta\rho_{max} = 1.15 \text{ e } \text{ Å}^{-3}$ $\Delta\rho_{min} = -1.10 \text{ e } \text{ Å}^{-3}$
264 parameters H atoms treated by a mixture of independent and constrained refinement	$\Delta \rho_{\rm min} = -1.10 \ {\rm e} \ {\rm A}^{-3}$

Table 1

Selected geometric parameters (Å, °).

Gd1-O2W	2.349 (4)	Gd1-O1	2.452 (4)
Gd1 - O1W	2.352 (4)	Gd1-O2	2.476 (5)
Gd1-O3	2.375 (4)	Gd1-O6 ⁱ	2.498 (4)
Gd1-O5 ⁱ	2.409 (4)	Gd1–O3 ⁱⁱ	2.643 (4)
Gd1-O4 ⁱⁱ	2.449 (4)		
O2W-Gd1-O1W	76.47 (17)	O5 ⁱ -Gd1-O2	129.05 (15)
O3-Gd1-O5 ⁱ	80.79 (15)	O4 ⁱⁱ -Gd1-O2	75.04 (15)
O3-Gd1-O4 ⁱⁱ	115.28 (14)	O1-Gd1-O2	52.51 (15)
O5 ⁱ -Gd1-O4 ⁱⁱ	82.01 (15)	O1-Gd1-O6 ⁱ	73.74 (16)
O3-Gd1-O1	152.25 (15)	O2-Gd1-O6 ⁱ	116.48 (15)
O5 ⁱ -Gd1-O1	79.66 (15)	O5 ⁱ -Gd1-O3 ⁱⁱ	74.25 (13)
O4 ⁱⁱ -Gd1-O1	81.18 (16)	O4 ⁱⁱ -Gd1-O3 ⁱⁱ	50.70 (13)
O3-Gd1-O2	150.15 (15)		

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

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

D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdots A$
0.847 (19)	1.86 (3)	2.688 (7)	166 (5)
0.849(19) 0.84(2)	1.94(3) 1.88(4)	2.773(6) 2.72(3)	167(7) 173(12)
0.84(2)	1.97 (5)	2.72 (3)	166 (12)
0.84(2)	2.39(7)	2.939(6)	123 (6)
	<i>D</i> -H 0.847 (19) 0.849 (19) 0.84 (2) 0.84 (2) 0.84 (2) 0.84 (2)	$\begin{array}{c ccc} D-H & H\cdots A \\ \hline 0.847 \ (19) & 1.86 \ (3) \\ 0.849 \ (19) & 1.94 \ (3) \\ 0.84 \ (2) & 1.88 \ (4) \\ 0.84 \ (2) & 1.97 \ (5) \\ 0.84 \ (2) & 2.39 \ (7) \\ 0.84 \ (2) & 2.2 \ (4) \\ 0.84 \ (4) & 2.8 \ (4) \ (4) \ $	$D-H$ $H\cdots A$ $D\cdots A$ 0.847 (19) 1.86 (3) 2.688 (7) 0.849 (19) 1.94 (3) 2.773 (6) 0.84 (2) 1.88 (4) 2.72 (3) 0.84 (2) 1.97 (5) 2.79 (3) 0.84 (2) 2.39 (7) 2.939 (6) 0.84 (2) 2.22 (4) 2.983 (6)

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

Water H atoms were located in a difference Fourier map and refined with distance restraints of O-H = 0.85(2) and $H \cdot \cdot \cdot H =$ 1.39 (1) Å. The remaining H atoms were placed in calculated positions in the riding-model approximation, with C-H = 0.93 Å and with $U_{iso}(H) = 1.2U_{eq}(C)$. The pyridine ring of the isonicotinate group is disordered; the occupancy was fixed as exactly 0.5. Distance restraints of C-C and C-N = 1.390 (5), and C···C and C···N = 2.780 (5) Å, were applied to the molecule. The highest peak is 1.15 e Å⁻³ at 0.91 Å from Gd1 and the deepest hole is -1.11 e Å⁻³ at 0.98 Å from Gd1.

Data collection: SMART (Bruker, 2001); cell refinement: SAINT (Bruker, 2001); data reduction: SAINT; program(s) used to solve structure: SHELXS97 (Sheldrick, 1997); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: SHELXTL (Bruker, 2001); software used to prepare material for publication: SHELXTL.

The authors thank the Liaoning University Programme for Excellent Talents for supporting this work (grant No. RC-05-11).

References

Bruker (2001). SAINT and SMART. Bruker AXS Inc., Madison, Wisconsin, USA.

Han, Z.-B., Li, X.-F., Cheng, X.-N. & Chen, X.-M. (2005). Z. Anorg. Allg. Chem. 631, 937–942.

Sheldrick, G. M. (1996). SADABS. University of Göttingen, Germany.

Sheldrick, G. M. (1997). SHELXS97 and SHELXL97. University of Göttingen, Germany.