

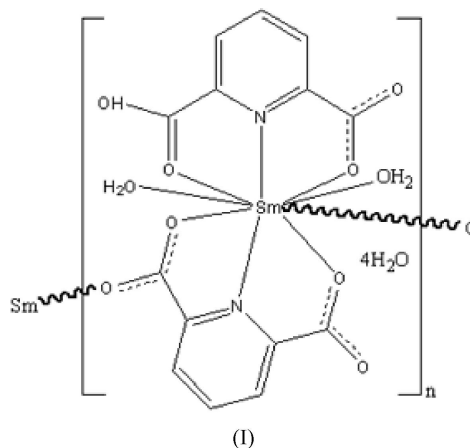
San-Hui Liu, Yi-Zhi Li\* and  
Qing-Jin MengCoordination Chemistry Institute, State Key  
Laboratory of Coordination Chemistry, Nanjing  
University, Nanjing, 210093, People's Republic  
of China

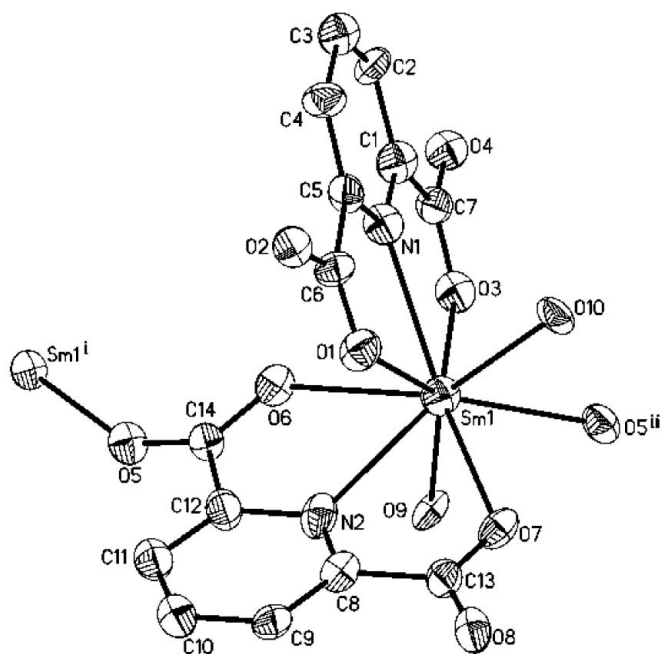
Correspondence e-mail: llyyz@nju.edu.cn

## Key indicators

Single-crystal X-ray study  
 $T = 293\text{ K}$   
Mean  $\sigma(\text{C}-\text{C}) = 0.012\text{ \AA}$   
 $R$  factor = 0.062  
 $wR$  factor = 0.156  
Data-to-parameter ratio = 13.7For details of how these key indicators were  
automatically derived from the article, see  
<http://journals.iucr.org/e>.**catena-Poly[[[diaqua(6-carboxypyridine-  
2-carboxylato- $\kappa^3\text{N},\text{O},\text{O}'$ )samarium(III)]-  
 $\mu$ -pyridine-2,6-dicarboxylato- $\kappa^4\text{N},\text{O},\text{O}':\text{O}''$ ]  
tetrahydrate]**In the title complex,  $[\text{Sm}(\text{C}_7\text{H}_3\text{O}_4)(\text{C}_7\text{H}_4\text{O}_4)(\text{H}_2\text{O})_2]\cdot 4\text{H}_2\text{O}$ , the coordination number of the Sm atom is nine. The pyridine-2,6-dicarboxylate ligand chelates to the Sm atom and bridges to the neighbouring Sm atoms to form infinite chains along the  $c$  axis.Received 7 April 2005  
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## Comment

In recent years, the use of lanthanide elements for constructing metal-organic framework (MOF) structures has attracted a lot of interest due to their high coordination numbers, along with special magnetic and luminescent properties (Benelli & Gatteschi, 2002; Ghosh & Bharadwaj, 2003). A search of the Cambridge Structural Database (February 2005 update; Allen, 2002) for pyridine-2,6-dicarboxylic acid ( $\text{pydcH}_2$ ) or the deprotonated ligand ( $\text{pydcH}^{-1}$  or  $\text{pydc}^{2-}$ ) coordinated to samarium yielded only five hits. The title compound, (I), which is a novel linear coordination polymer of samarium with pyridine-2,6-dicarboxylate ligands, is reported here.The asymmetric unit of (I) is composed of one  $\text{Sm}^{\text{III}}$  ion, one  $\text{PydcH}^{-1}$  and one  $\text{Pydc}^{2-}$  ligand, two coordinated water molecules and four water molecules of crystallization. Atom Sm1 is coordinated by  $\text{pydcH}^{-}$  (atoms O1, N1 and O3),  $\text{pydc}^{2-}$  (O6, N2 and O7), and two aqua ligands (O9 and O10); the Sm atom is also bonded to one bridging atom, O5<sup>iii</sup> [symmetry code: (iii)  $x, y, z - 1$ ], from a neighbouring  $\text{pydc}^{2-}$  ligand, giving an overall nine-coordination (Fig. 1). The  $\text{pydc}^{2-}$  group chelates to the Sm1 atom and bridges to the neighbouring Sm1<sup>i</sup> atom [symmetry code: (i)  $x, \frac{3}{2} - y, z + \frac{1}{2}$ ], forming an infinite chain along the  $c$  axis (Fig. 2). Two neighbouring chains are linked to form a ladder-like band through O—H $\cdots$ O hydrogen bonds (Table 2 and Fig. 3). The ladder-like


**Figure 1**

The structure of (I), with displacement ellipsoids drawn at the 50% probability level. All H atoms and uncoordinated water molecules have been omitted for clarity. [Symmetry codes: (i)  $x, \frac{3}{2} - y, z + \frac{1}{2}$ ; (ii)  $x, \frac{3}{2} - y, z - \frac{1}{2}$ ;

bands form a layer structure parallel to (100) by hydrogen bonds with the uncoordinated water molecules, and adjacent layers are linked together through hydrogen bonds, resulting in a three-dimensional framework.

## Experimental

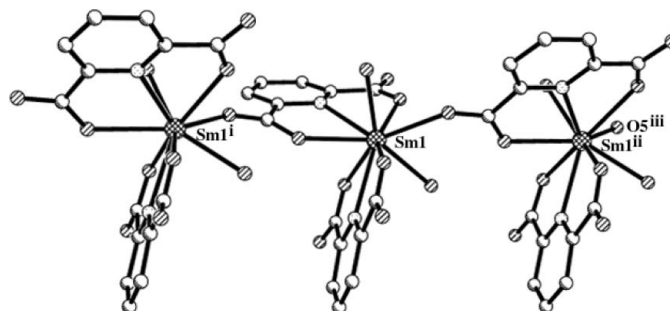
$\text{Sm}_2\text{O}_3$  was acquired from Aldrich, and pyridine-2,6-dicarboxylic acid was synthesized according to the literature (Singer & McElvain, 1935). To a solution of pyridine-2,6-dicarboxylic acid (167 mg, 1 mmol) in water (60 ml) and sodium hydroxide (1 M, 2 ml),  $\text{Sm}_2\text{O}_3$  (349 mg, 1 mmol) was added. The mixture was stirred at 373 K for 12 h and then filtered. Single crystals of (I) suitable for X-ray analysis were grown from the filtrate after six weeks (yield 65%). Compound (I) is stable in air and soluble in water.

### Crystal data

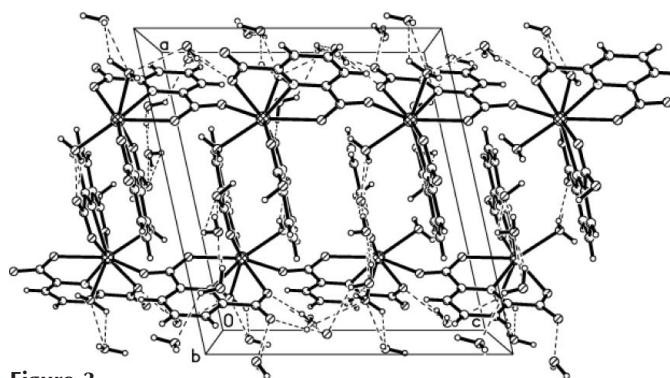
$\text{C}_{14}\text{H}_{11}\text{N}_2\text{O}_{10}\text{Sm}\cdot 4\text{H}_2\text{O}$   
 $M_r = 589.66$   
 Monoclinic,  $P2_1/c$   
 $a = 13.973$  (3) Å  
 $b = 11.203$  (2) Å  
 $c = 12.830$  (3) Å  
 $\beta = 102.367$  (4)°  
 $V = 1961.8$  (7) Å<sup>3</sup>  
 $Z = 4$   
 $D_x = 1.996$  Mg m<sup>-3</sup>  
 Mo  $K\alpha$  radiation  
 Cell parameters from 935 reflections  
 $\theta = 2.4\text{--}25.6^\circ$   
 $\mu = 3.07$  mm<sup>-1</sup>  
 $T = 293$  (2) K  
 Prism, colourless  
 $0.20 \times 0.14 \times 0.12$  mm

### Data collection

Bruker SMART APEX CCD area-detector diffractometer  
 $\varphi$  and  $\omega$  scans  
 Absorption correction: multi-scan (SADABS; Bruker, 2000)  
 $T_{\min} = 0.61, T_{\max} = 0.69$   
 10 285 measured reflections  
 3846 independent reflections  
 2849 reflections with  $I > 2\sigma(I)$   
 $R_{\text{int}} = 0.044$   
 $\theta_{\text{max}} = 26.0^\circ$   
 $h = -16 \rightarrow 17$   
 $k = -13 \rightarrow 13$   
 $l = -13 \rightarrow 15$


**Figure 2**

The infinite chain along the  $c$  axis. [Symmetry codes: (i)  $x, \frac{3}{2} - y, z + \frac{1}{2}$ ; (ii)  $x, \frac{3}{2} - y, z - \frac{1}{2}$ ; (iii)  $x, y, z - 1$ .]


**Figure 3**

The crystal packing of (I), viewed down the  $b$  axis. Dashed lines indicate hydrogen bonds.

### Refinement

Refinement on  $F^2$   
 $R[F^2 > 2\sigma(F^2)] = 0.062$   
 $wR(F^2) = 0.156$   
 $S = 1.06$   
 3846 reflections  
 280 parameters  
 H-atom parameters constrained

$$w = 1/[\sigma^2(F_o^2) + (0.09P)^2 + 1.99P]$$

where  $P = (F_o^2 + 2F_c^2)/3$   
 $(\Delta/\sigma)_{\text{max}} < 0.001$   
 $\Delta\rho_{\text{max}} = 0.98 \text{ e \AA}^{-3}$   
 $\Delta\rho_{\text{min}} = -2.01 \text{ e \AA}^{-3}$

**Table 1**

Selected geometric parameters (Å, °).

Sm1—O9	2.388 (5)	Sm1—O6	2.524 (6)
Sm1—O10	2.423 (6)	Sm1—N2	2.537 (8)
Sm1—O3	2.426 (6)	Sm1—O1	2.537 (6)
Sm1—O5 <sup>i</sup>	2.446 (6)	Sm1—N1	2.556 (7)
Sm1—O7	2.476 (6)		
O9—Sm1—O10	141.5 (2)	O5 <sup>i</sup> —Sm1—N2	128.9 (2)
O9—Sm1—O3	79.90 (19)	O7—Sm1—N2	62.3 (2)
O10—Sm1—O3	97.26 (19)	O6—Sm1—N2	62.5 (2)
O9—Sm1—O5 <sup>i</sup>	71.6 (2)	O9—Sm1—O1	141.0 (2)
O10—Sm1—O5 <sup>i</sup>	70.7 (2)	O10—Sm1—O1	71.2 (2)
O3—Sm1—O5 <sup>i</sup>	74.65 (19)	O3—Sm1—O1	124.7 (2)
O9—Sm1—O7	86.04 (18)	O5 <sup>i</sup> —Sm1—O1	139.1 (2)
O10—Sm1—O7	78.56 (19)	O7—Sm1—O1	81.21 (18)
O3—Sm1—O7	151.38 (19)	O6—Sm1—O1	84.23 (19)
O5 <sup>i</sup> —Sm1—O7	77.3 (2)	N2—Sm1—O1	65.8 (2)
O9—Sm1—O6	72.88 (19)	O9—Sm1—N1	133.5 (2)
O10—Sm1—O6	143.9 (2)	O10—Sm1—N1	73.3 (2)
O3—Sm1—O6	75.08 (19)	O3—Sm1—N1	62.4 (2)
O5 <sup>i</sup> —Sm1—O6	136.45 (19)	O5 <sup>i</sup> —Sm1—N1	118.5 (2)
O7—Sm1—O6	124.2 (2)	O7—Sm1—N1	139.4 (2)
O9—Sm1—N2	75.5 (2)	O6—Sm1—N1	72.0 (2)
O10—Sm1—N2	124.8 (2)	N2—Sm1—N1	112.5 (2)
O3—Sm1—N2	135.5 (2)	O1—Sm1—N1	62.4 (2)

Symmetry code: (i)  $x, \frac{3}{2} - y, z - \frac{1}{2}$ .

**Table 2**  
Hydrogen-bonding geometry (Å, °).

<i>D</i> —H... <i>A</i>	<i>D</i> —H	H... <i>A</i>	<i>D</i> ... <i>A</i>	<i>D</i> —H... <i>A</i>
O2—H2A...O3 <sup>ii</sup>	0.96	2.60	3.515 (9)	161
O9—H9A...O12	0.96	2.50	3.295 (8)	140
O9—H9C...O14 <sup>iii</sup>	0.96	2.03	2.720 (7)	127
O10—H10A...O6 <sup>i</sup>	0.85	2.13	2.707 (8)	125
O10—H10C...O1	0.85	2.48	2.889 (8)	111
O10—H10C...O4 <sup>ii</sup>	0.85	2.01	2.720 (9)	141
O11—H11B...O7	0.85	2.08	2.912 (8)	164
O12—H12A...O14 <sup>iv</sup>	0.85	2.39	3.004 (9)	130
O12—H12B...O3	0.85	2.52	2.945 (8)	112
O12—H12B...O13	0.85	2.07	2.775 (8)	140
O13—H13B...O4	0.85	1.87	2.642 (8)	150
O14—H14B...O8 <sup>v</sup>	0.95	1.73	2.683 (8)	176
O14—H14C...O11 <sup>v</sup>	0.86	2.17	2.895 (8)	142

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

O-bound H atoms were located from difference density maps (O—H = 0.85–0.96 Å) and C-bound H atoms were positioned geome-

trically (C—H = 0.93 Å). All H atoms were refined as riding, with isotropic displacement parameters 1.2 times the  $U_{eq}$  value of the parent atom. The maximum and minimum electron-density peaks are located 1.17 and 0.55 Å, respectively, from atom Sm1.

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

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## References

- Allen, F. H. (2002). *Acta Cryst.* **B58**, 380–388.  
 Benelli, C. & Gatteschi, D. (2002). *Chem. Rev.* **102**, 2369–2387.  
 Bruker (2000). *SMART* (Version 5.625), *SAINTE* (Version 6.01), *SHELXTL* (Version 6.10) and *SADABS* (Version 2.03). Bruker AXS Inc., Madison, Wisconsin, USA.  
 Ghosh, S. K. & Bharadwaj, P. K. (2003). *Inorg. Chem.* **42**, 8250–8254.  
 Singer, A. W. & McElvain, S. M. (1935). *J. Am. Chem. Soc.* **57**, 1135–1137.