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Key indicators

Single-crystal X-ray study  
 T = 298 K  
 Mean  $\sigma(\text{C}-\text{C}) = 0.007 \text{ \AA}$   
 H-atom completeness 86%  
 Disorder in solvent or counterion  
 R factor = 0.032  
 wR factor = 0.071  
 Data-to-parameter ratio = 14.9

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

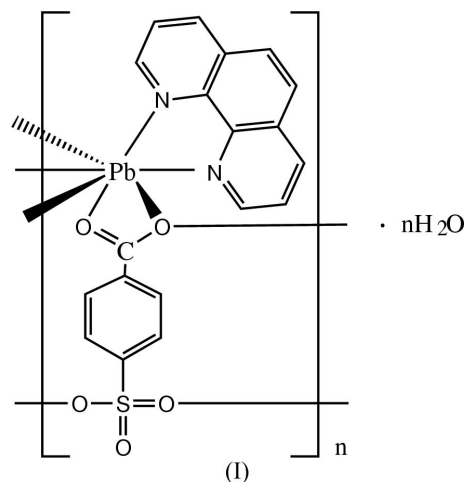
Poly[(1,10-phenanthroline- $\kappa^2N,N'$ )lead(II)  
 $\mu_4$ -(4-sulfonatobenzoato- $\kappa^4O,O':O'',O'''$   
 hydrate)]

In the title polymeric compound,  $[\text{Pb}(\text{C}_{12}\text{H}_8\text{N}_2)(\text{C}_7\text{H}_4\text{O}_5\text{S})]_n \cdot n\text{H}_2\text{O}$ , the seven-coordinate  $\text{Pb}^{\text{II}}$  atom is surrounded by one 1,10-phenanthroline (phen) and four 4-sulfonatobenzoate (sb) ligands in a hemidirected coordination geometry. Both the sulfonyl and carboxy groups of the sb ligand bridge the neighboring  $\text{Pb}^{\text{II}}$  atoms to form a two-dimensional sheet structure.

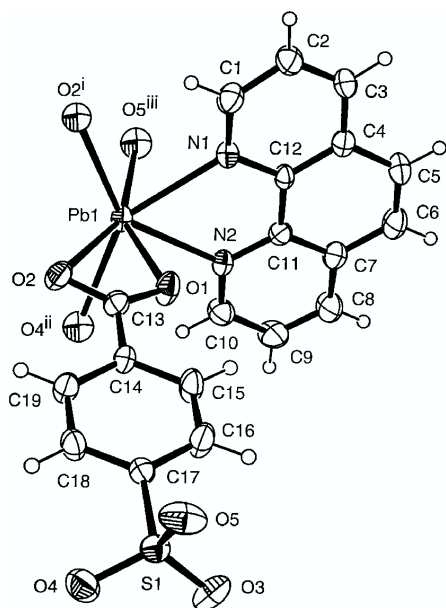
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Comment

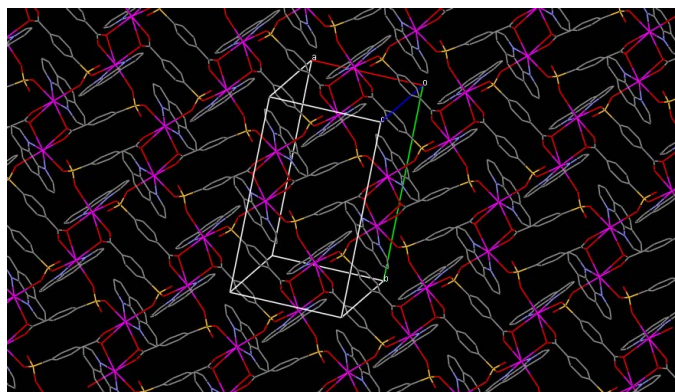
The sulfonate group exhibits a different coordination mode compared to the carboxylate group (Zheng *et al.*, 2003; Wang *et al.*, 2004; Starynowicz, 2000). 4-Sulfonatobenzoate (sb), which combines sulfonate and carboxylate groups, may be used as a bridging ligand to construct novel complex architectures. A search of the Cambridge Structure Database (November 2004 update; Allen, 2002) shows that the structures of five sb metal complexes have been reported, the sulfonate group adopting a monodentate mode (Yuan *et al.*, 2001; Fan, Xiao, Zhang & Zhu, L.-G., 2004; Fan, Xiao, Zhang, Cai & Zhu, 2004) or a *syn-syn* bridge mode (Xiong *et al.*, 2001) in these complexes. We report here the crystal structure of the title polymeric  $\text{Pb}^{\text{II}}$  complex, (I), which exhibits a different coordination mode for the sulfonate group from those reported previously.



The structure of a segment of (I) is presented in Fig. 1 to show the coordination geometry around the  $\text{Pb}^{\text{II}}$  atom. Each seven-coordinate  $\text{Pb}^{\text{II}}$  atom is surrounded by one 1,10-phenanthroline (phen) and four sb ligands. While the carboxylate group of the sb ligand chelates a  $\text{Pb}^{\text{II}}$  atom, the O2 atom of the carboxylate group coordinates to a neighboring  $\text{Pb}^{\text{II}}$  atom to form a centrosymmetric  $\text{Pb}_2\text{O}_2$  four-membered ring, with a  $\text{Pb} \cdots \text{Pb}$  separation of 4.5813 (4)  $\text{ \AA}$ . The bridging


**Figure 1**

The structure of a segment of (I), showing the coordination geometry. Displacement ellipsoids are drawn at the 40% probability level. Uncoordinated water molecules are omitted for clarity. Symmetry codes (i):  $-x, 1-y, 1-z$ ; (ii):  $1-x, 1-y, 1-z$ ; (iii):  $\frac{1}{2}-x, -\frac{1}{2}+y, z$ .


**Figure 2**

A view of the two-dimensional network of (I). The uncoordinated water molecules and H atoms are omitted for clarity.

mode of the carboxylate is *anti-skew*, with a dihedral angle of  $10.1(8)^\circ$  between the carboxyl group and the benzene ring. The sulfonate group of the sb ligand links two neighboring Pb<sup>II</sup> atoms in a skew-skew bridging mode, the Pb<sup>II</sup>–Pb separation being  $7.1712(4)$  Å. Each sb ligand, therefore, bridges four Pb<sup>II</sup> atoms to form a two-dimensional sheet network (Fig. 2). Three kinds of metal-containing rings, [Pb<sup>II</sup>–O–Pb<sup>II</sup>–O], [Pb<sup>II</sup>–sb–Pb<sup>II</sup>–SO<sub>3</sub>] and [Pb<sup>II</sup>–sb–Pb<sup>II</sup>–sb], are observed in the sheet network. Within the [Pb<sup>II</sup>–sb–Pb<sup>II</sup>–sb] moiety, parallel mean planes of two sb ligands are separated from each other by a face-to-face distance of  $3.57(3)$  Å.

The coordination geometry around the Pb<sup>II</sup> atom in (I) may be described as ‘hemidirected’ (Shimoni-Livny *et al.*, 1998), a significant gap being located in the position *trans* to the phen ligand. In the empty direction, some short contacts are

observed, *viz.* Pb<sup>II</sup>–O6B<sup>ii</sup> =  $3.34(2)$  Å and Pb<sup>II</sup>–O3<sup>ii</sup> =  $3.828(4)$  Å [symmetry code (ii):  $1-x, 1-y, 1-z$ ]. This is similar to the situation found in [Pb(DOTAM)](ClO<sub>4</sub>)<sub>2</sub>·5H<sub>2</sub>O (DOTAM = 1,4,7,10-tetrakis(carbamoylmethyl)-1,4,7,10-tetraazacyclododecane) (Hancock *et al.*, 2004).

## Experimental

A mixture of Pb(NO<sub>3</sub>)<sub>2</sub> (0.084 g, 0.25 mmol), KHsb (0.061 g, 0.25 mmol), phen (0.046 g, 0.23 mmol) and H<sub>2</sub>O (15 ml) was placed in a 30 ml stainless steel reactor with a Teflon liner and heated at 423 K for 24 h. After cooling, colorless block-shaped crystals of (I) were obtained. Elemental and TG analyses clearly indicate that the compound contains a water molecule, with a formula of [Pb(C<sub>12</sub>H<sub>8</sub>N<sub>2</sub>)(C<sub>7</sub>H<sub>4</sub>O<sub>5</sub>S)]·H<sub>2</sub>O. Analysis calculated for C<sub>19</sub>H<sub>14</sub>N<sub>2</sub>O<sub>6</sub>·PbS: C 37.68, H 2.33, N 4.63%; found: C 37.42, H 2.34, N 4.46%. Weight loss in the temperature range 363–408 K corresponds to the release of the water molecule (calculated 2.98%, found 3.07%).

### Crystal data

[Pb(C<sub>12</sub>H<sub>8</sub>N<sub>2</sub>)(C<sub>7</sub>H<sub>4</sub>O<sub>5</sub>S)]·H<sub>2</sub>O  
 $M_r = 605.57$   
 Orthorhombic, *Pbca*  
 $a = 9.8745(7)$  Å  
 $b = 17.3467(11)$  Å  
 $c = 21.7544(14)$  Å  
 $V = 3726.3(4)$  Å<sup>3</sup>  
 $Z = 8$   
 $D_x = 2.159$  Mg m<sup>-3</sup>

Mo K $\alpha$  radiation  
 Cell parameters from 3784 reflections  
 $\theta = 2.5$ – $25.2^\circ$   
 $\mu = 9.21$  mm<sup>-1</sup>  
 $T = 298(2)$  K  
 Block, colorless  
 $0.29 \times 0.14 \times 0.07$  mm

### Data collection

Bruker SMART area-detector diffractometer  
 $\varphi$  and  $\omega$  scans  
 Absorption correction: multi-scan (SADABS; Bruker, 2002)  
 $T_{\min} = 0.206$ ,  $T_{\max} = 0.523$   
 21368 measured reflections

4064 independent reflections  
 3160 reflections with  $I > 2\sigma(I)$   
 $R_{\text{int}} = 0.048$   
 $\theta_{\max} = 27.0^\circ$   
 $h = -12 \rightarrow 12$   
 $k = -22 \rightarrow 18$   
 $l = -25 \rightarrow 27$

### Refinement

Refinement on  $F^2$   
 $R[F^2 > 2\sigma(F^2)] = 0.032$   
 $wR(F^2) = 0.071$   
 $S = 1.01$   
 4064 reflections  
 273 parameters  
 H-atom parameters constrained

$w = 1/[\sigma^2(F_o^2) + (0.0278P)^2 + 2.5851P]$   
 where  $P = (F_o^2 + 2F_c^2)/3$   
 $(\Delta/\sigma)_{\max} = 0.001$   
 $\Delta\rho_{\max} = 1.32$  e Å<sup>-3</sup>  
 $\Delta\rho_{\min} = -0.75$  e Å<sup>-3</sup>

**Table 1**

Selected geometric parameters (Å, °).

Pb1–O1	2.439 (3)	Pb1–N1	2.546 (4)
Pb1–O2	2.888 (3)	Pb1–N2	2.540 (4)
Pb1–O2 <sup>i</sup>	2.715 (3)	S1–O3	1.441 (4)
Pb1–O4 <sup>ii</sup>	2.699 (4)	S1–O4	1.441 (4)
Pb1–O5 <sup>iii</sup>	2.694 (4)	S1–O5	1.446 (4)
O1–Pb1–O2	48.28 (10)	O5 <sup>iii</sup> –Pb1–O2 <sup>i</sup>	94.01 (11)
O1–Pb1–O2 <sup>i</sup>	96.05 (11)	N1–Pb1–O2	113.20 (11)
O1–Pb1–O4 <sup>ii</sup>	87.35 (12)	N1–Pb1–O2 <sup>i</sup>	77.50 (11)
O1–Pb1–O5 <sup>iii</sup>	151.95 (13)	N1–Pb1–O4 <sup>ii</sup>	142.56 (12)
O1–Pb1–N1	80.65 (12)	N1–Pb1–O5 <sup>iii</sup>	76.08 (12)
O1–Pb1–N2	78.68 (12)	N2–Pb1–N1	65.22 (13)
O2 <sup>i</sup> –Pb1–O2	70.32 (12)	N2–Pb1–O2	124.04 (11)
O4 <sup>ii</sup> –Pb1–O2	82.72 (10)	N2–Pb1–O2 <sup>i</sup>	142.72 (11)
O4 <sup>ii</sup> –Pb1–O2 <sup>i</sup>	139.34 (11)	N2–Pb1–O4 <sup>ii</sup>	77.73 (12)
O5 <sup>iii</sup> –Pb1–O2	158.36 (11)	N2–Pb1–O5 <sup>iii</sup>	77.49 (12)
O5 <sup>iii</sup> –Pb1–O4 <sup>ii</sup>	101.74 (11)		

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

The aromatic H atoms were positioned geometrically and refined using the riding-model approximation [ $C-H = 0.93 \text{ \AA}$ ,  $U_{iso}(H) = 1.2U_{eq}(C)$ ]. The uncoordinated water is disordered over two sites; occupancies were initially refined and converged to 0.532 (3) and 0.468 (3), respectively, and were then fixed at 0.53 (major component) and 0.47 (minor component) in the final cycles of refinement. H atoms of the disordered water molecule were not located. The highest peak in the final difference Fourier map lies  $0.92 \text{ \AA}$  from atom Pb.

Data collection: *SMART* (Bruker, 2002); cell refinement: *SAINTE* (Bruker, 2002); data reduction: *SAINTE*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *ORTEP-3 for Windows* (Farrugia, 1997); software used to prepare material for publication: *WinGX* (Farrugia, 1999).

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