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Key indicators

Single-crystal X-ray study T = 298 KMean σ (C–C) = 0.012 Å R factor = 0.034 wR factor = 0.083 Data-to-parameter ratio = 11.4

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

Poly[[tetraaquabis(µ₅-1-carboxybenzene-3,5-dicarboxylato)dilead(II)] monohydrate]

Crystals of the title compound, $\{[Pb_2(C_9H_4O_6)_2(H_2O)_4]$ - $H_2O\}_n$, were obtained from an aqueous solution. In the crystal structure, each Pb^{II} cation is nine-coordinated by seven O atoms from five carboxylate groups of five 1-carboxybenzene-3,5-dicarboxylate (TBC) dianions and two water molecules. Pairs of Pb atoms are bridged by two tridentate carboxylate groups of two TBC dianions, yielding centrosymmetric dinuclear complexes. These are connected by TBC dianions, forming a three-dimensional network.

Comment

Lead(II) is capable of accommodating different coordination numbers and geometries with or without a stereochemically active lone pair (Wang & Vittal, 2003). Of these, a number of polymeric Pb^{II} compounds have been structurally characterized (Cecconi et al., 2003; Bridgewater & Parkin, 2000; Ying et al., 2003). The absence of crystal field stabilization energy effects also allows Pb^{II} cations to adopt a range of different coordination geometries not restricted to octahedral, tetrahedral or square planar (Foreman et al., 2000). The benzene-1,3,5-tricarboxylic acid (TBCA) ligand binds to metal atoms in a variety of bonding modes. When TBCA binds to lead ions, various coordination modes and conformations can be obtained, making this ligand particularly useful for the study of the coordination chemistry of Pb²⁺. Thus, we have selected the Pb-TBCA system to extend this research and we present here the crystal structure of the title compound, (I).



Compound (I) crystallizes with one Pb^{2+} and one TBC dianion in the asymmetric unit (Fig. 1). The Pb atom has a coordination number of nine. Each Pb ion is coordinated by seven O atoms from five carboxylate groups from five TBC anions. Two O atoms from water molecules complete the

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The contents of the asymmetric unit of (I), together with symmetryequivalent atoms to complete the coordination of Pb1. Symmetry codes as in Table 1.





Coordination mode of the TBC dianion. Symmetry operations as in Table 1, plus (v) x, y, 1 + z; (vi) 1 + x, 1 + y, 1 + z.

coordination environment. The coordination geometry about Pb cannot be derived from a simple polyhedron. The geometry of the TBC dianion is unexceptional, and two of its three carboxylate groups are deprotonated. Each dianion coordinates to five Pb^{2+} ; Fig. 2 shows the coordination mode. One of the deprotonated carboxylate groups coordinates to three Pb: O1 and O2 chelate a Pb cation, and both O1 and O2 coordinate to one Pb cation. The second carboxylate group (O3 and O4) coordinates to one Pb. The protonated carboxylate group is monodentate. Hence, pairs of Pb^{II} ions are bridged by two tridentate carboxylate groups, forming centrosymmetric dinuclear units with the inversion centre halfway between the



Figure 3 Chains derived from edge-sharing PbO₉ polyhedra.

two Pb. These units form polymeric chains that run along the a axis (Fig. 3). The chains are linked together by the TBC anions, yielding a three-dimensional framework.

There are also $O-H \cdots O$ hydrogen bonds (Table 2), which connect the coordination polymer and water molecules and help to stabilize the crystal structure.

Experimental

The title compound was synthesized by adding a solution of benzene-1,3,5-tricarboxylic acid (0.04 g, 0.2 mmol) and 2,2'-dithiosalicylic acid (0.06 g, 0.2 mmol) in dimethylformamide (10 ml) dropwise to a stirred solution of lead nitrate (0.07 g, 0.2 mmol) in water (10 ml) at 298 K. The reaction mixture was filtered and the filtrate allowed to stand for about six weeks until colourless prismatic crystals of (I) were obtained. Crystals suitable for X-ray diffraction were collected by filtration, washed with water and ethanol and dried in air.

Crystal data

| $Pb_2(C_9H_4O_6)_2(H_2O)_4] \cdot H_2O$ | $V = 557.91 (10) \text{ Å}^3$ |
|---|---|
| $M_r = 910.64$ | Z = 1 |
| Triclinic, P1 | $D_x = 2.710 \text{ Mg m}^{-3}$ |
| a = 7.4106 (6) Å | Mo $K\alpha$ radiation |
| b = 8.2178 (7) Å | $\mu = 15.16 \text{ mm}^{-1}$ |
| c = 10.1424 (9) Å | T = 298 (2) K |
| $\alpha = 94.694 \ (10)^{\circ}$ | Prism, colourless |
| $\beta = 105.165 \ (10)^{\circ}$ | $0.10 \times 0.06 \times 0.05 \text{ mm}$ |
| $\gamma = 108.024 \ (9)^{\circ}$ | |
| | |

Data collection

Bruker APEX area-detector3941 mdiffractometer1949 in ω scans1896 mAbsorption correction: numerical $R_{int} = 0.348$; Bruker, 2002) $T_{min} = 0.344$, $T_{max} = 0.465$ $\theta_{max} = 0.465$

Refinement

Refinement on F^2 $R[F^2 > 2\sigma(F^2)] = 0.034$ $wR(F^2) = 0.083$ S = 1.29S = 1.29(1949 reflections)171 parameters(1949 reflections)H-atom parameters constrained(1940 reflections)

3941 measured reflections 1949 independent reflections 1896 reflections with $I > 2\sigma(I)$ $R_{\text{int}} = 0.030$ $\theta_{\text{max}} = 25.1^{\circ}$

$$\begin{split} &w = 1/[\sigma^2(F_{\rm o}^2) + (0.017P)^2 \\ &+ 3.5615P] \\ &where \ P = (F_{\rm o}^2 + 2F_{\rm c}^2)/3 \\ (\Delta/\sigma)_{\rm max} < 0.001 \\ \Delta\rho_{\rm max} = 1.48 \ {\rm e} \ {\rm \AA}^{-3} \\ \Delta\rho_{\rm min} = -1.57 \ {\rm e} \ {\rm \AA}^{-3} \\ {\rm Extinction \ correction: \ SHELXL97} \\ {\rm Extinction \ coefficient: \ 0.0066 \ (8)} \end{split}$$

Table 1Selected bond lengths (Å).

| Pb1-O4 ⁱ | 2.562 (6) | Pb1-O1 ⁱⁱ | 2.745 (6) | |
|---------------------|-----------|-----------------------|-----------|--|
| Pb1-O1 | 2.617 (6) | Pb1-O5 ⁱⁱⁱ | 2.782 (6) | |
| Pb1-O2 | 2.641 (6) | $Pb1-O2^{iv}$ | 2.815 (6) | |
| Pb1-O7 | 2.652 (9) | Pb1-O8 | 2.875 (8) | |
| $Pb1 - O3^{i}$ | 2 702 (6) | | | |

Symmetry codes: (i) x, y, z - 1; (ii) -x + 2, -y + 1, -z; (iii) x - 1, y - 1, z - 1; (iv) -x + 1, -y + 1, -z.

Table 2

Hydrogen-bond geometry (Å, °).

| $O6-H6\cdots O3^{v}$ 0.82 1.83 2.629 (8) 164 | $D - H \cdots A$ | D-H | $H \cdot \cdot \cdot A$ | $D \cdot \cdot \cdot A$ | $D - H \cdot \cdot \cdot A$ |
|--|----------------------|------|-------------------------|-------------------------|-----------------------------|
| | $O6-H6\cdots O3^{v}$ | 0.82 | 1.83 | 2.629 (8) | 164 |

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

H atoms were positioned geometrically and allowed to ride on their parent atoms at distances of C-H = 0.93 Å, with $U_{iso}(H) =$ $1.2U_{eq}(C)$, and O-H = 0.82 Å, with $U_{iso}(H) = 1.5U_{eq}(O)$. Water H atoms appear to be disordered and were not considered in the structure model. The highest peak and deepest hole of the residual electron density are 1.06 and 0.92 Å, respectively, from Pb1. Data collection: *SMART* (Bruker, 2002); cell refinement: *SAINT* (Bruker, 2002); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *XP* in *SHELXTL* (Bruker, 2002); software used to prepare material for publication: *SHELXL97*.

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