

Shan Gao<sup>a</sup> and Seik Weng Ng<sup>b\*</sup><sup>a</sup>College of Chemistry and Materials Science, Heilongjiang University, Harbin 150080, People's Republic of China, and <sup>b</sup>Department of Chemistry, University of Malaya, Kuala Lumpur 50603, Malaysia

Correspondence e-mail: seikweng@um.edu.my

## Key indicators

Single-crystal X-ray study  
 $T = 295\text{ K}$   
Mean  $\sigma(\text{C}-\text{C}) = 0.003\text{ \AA}$   
H-atom completeness 82%  
Disorder in solvent or counterion  
 $R$  factor = 0.045  
 $wR$  factor = 0.131  
Data-to-parameter ratio = 16.9For details of how these key indicators were automatically derived from the article, see <http://journals.iucr.org/e>.*catena*-Poly[[[diaquacalcium(II)]-di- $\mu$ -3-carboxyphenoxyacetato] dihydrate]

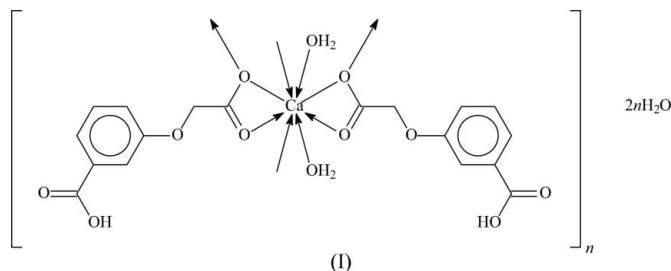
The Ca atom in the title compound,  $\{[\text{Ca}(\text{HO}_2\text{CC}_6\text{H}_4\text{OCH}_2\text{CO}_2)_2(\text{H}_2\text{O})_2]\cdot 2\text{H}_2\text{O}\}_n$ , is eight-coordinate. The  $\text{HO}_2\text{CC}_6\text{H}_4\text{OCH}_2\text{CO}_2^-$  monoanion binds in a chelating mode; it also functions as a bridge, linking adjacent Ca atoms into a zigzag chain. Hydrogen bonds link the chains into a three-dimensional network. The Ca atom lies on a special position of site symmetry 2.

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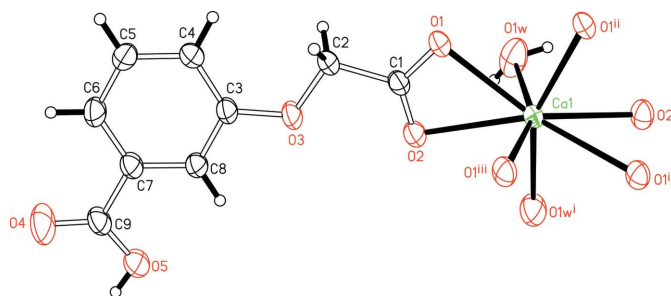
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## Comment

A preceding study (Gao & Ng, 2006) reports the structure of the cadmium derivative of 3-carboxyphenoxyacetic acid; the acid is doubly deprotonated and the dianion links the water-coordinated Cd atoms into a linear chain.



With calcium in place of cadmium, a similar synthesis, but employing less than half the stoichiometric quantity of the acid, has ensured that the acid is only mono-deprotonated. The calcium derivative is formally the water-coordinated complex,  $(\text{H}_2\text{O})_2(\text{HO}_2\text{CC}_6\text{H}_4\text{OCH}_2\text{CO}_2)_2\text{Ca}$ , which crystallizes as a dihydrate (Scheme, Fig. 1). The acid H atom is retained by the carboxyl unit that is directly connected to the benzene ring, and the carboxylic acid portion engages in hydrogen-bonding interactions (Table 2), giving rise to a three-dimensional network.



**Figure 1**  
ORTEP plot of a portion of the chain structure of  $\text{Ca}(\text{C}_9\text{H}_7\text{O}_5)_2(\text{H}_2\text{O})_2\cdot 2\text{H}_2\text{O}$ . Displacement ellipsoids are drawn at the 50% probability level, and H atoms are drawn as spheres of arbitrary radii. The disordered uncoordinated water molecule is not shown.

The carboxylate group of the oxyacetate arm binds in a chelating mode; additionally, one of the two O atoms interacts with an adjacent Ca atom. Such a bridging interaction leads to the formation of a zigzag chain that runs along the *c* axis (Fig. 2). The Ca atom is eight-coordinate; however, the geometry cannot be ascribed to a regular polyhedron.

**Experimental**

Calcium nitrate tetrahydrate (0.78 g, 5 mmol) and 3-carboxyphenoxyacetic acid (0.39 g, 2 mmol) were dissolved in a small volume of hot water. Colorless prisms separated from the solution after several days. C&H analysis. Calc. for C<sub>18</sub>H<sub>22</sub>O<sub>14</sub>Ca: C 43.02, H 4.42%. Found: C 43.06, H 4.45%.

*Crystal data*

[Ca(C<sub>9</sub>H<sub>7</sub>O<sub>5</sub>)<sub>2</sub>(H<sub>2</sub>O)<sub>2</sub>].2H<sub>2</sub>O  
*M<sub>r</sub>* = 502.44  
 Monoclinic, *C2/c*  
*a* = 26.902 (5) Å  
*b* = 14.492 (3) Å  
*c* = 6.057 (1) Å  
 $\beta$  = 101.28 (3)°  
*V* = 2315.6 (8) Å<sup>3</sup>  
*Z* = 4  
*D<sub>x</sub>* = 1.441 Mg m<sup>-3</sup>  
 Mo *K*α radiation  
 Cell parameters from 10195 reflections  
 $\theta$  = 3.1–27.4°  
 $\mu$  = 0.34 mm<sup>-1</sup>  
*T* = 295 (2) K  
 Prism, colorless  
 0.36 × 0.25 × 0.18 mm

*Data collection*

Rigaku RAXIS-RAPID IP diffractometer  
 $\omega$  scans  
 Absorption correction: multi-scan *ABSCOR* (Higashi, 1995)  
*T<sub>min</sub>* = 0.691, *T<sub>max</sub>* = 0.941  
 10847 measured reflections  
 2622 independent reflections  
 2340 reflections with *I* > 2σ(*I*)  
*R<sub>int</sub>* = 0.021  
 $\theta_{max}$  = 27.4°  
*h* = -34 → 33  
*k* = -18 → 18  
*l* = -7 → 7

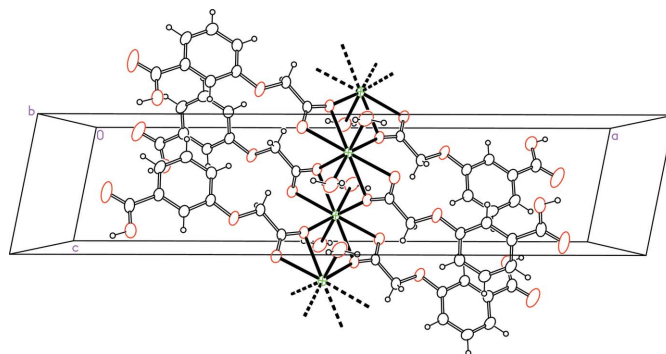
*Refinement*

Refinement on *F*<sup>2</sup>  
*R* [*F*<sup>2</sup> > 2σ(*F*<sup>2</sup>)] = 0.045  
*wR* (*F*<sup>2</sup>) = 0.131  
*S* = 1.04  
 2622 reflections  
 155 parameters  
 H-atom parameters constrained  
 $w = 1/[\sigma^2(F_o^2) + (0.0777P)^2 + 2.6608P]$   
 where  $P = (F_o^2 + 2F_c^2)/3$   
 $(\Delta/\sigma)_{max} = 0.001$   
 $\Delta\rho_{max} = 0.42 \text{ e \AA}^{-3}$   
 $\Delta\rho_{min} = -0.37 \text{ e \AA}^{-3}$

**Table 1**  
 Selected geometric parameters (Å, °).

Ca1—O1	2.488 (2)	Ca1—O2	2.537 (2)
Ca1—O1 <sup>i</sup>	2.488 (2)	Ca1—O2 <sup>i</sup>	2.537 (2)
Ca1—O1 <sup>ii</sup>	2.386 (1)	Ca1—O1 <sub>w</sub>	2.393 (2)
Ca1—O1 <sup>iii</sup>	2.386 (1)	Ca1—O1 <sub>w</sub> <sup>i</sup>	2.393 (2)
O1—Ca1—O1 <sup>i</sup>	146.2 (1)	O1 <sup>ii</sup> —Ca1—O2	126.3 (1)
O1—Ca1—O1 <sup>ii</sup>	74.4 (1)	O1 <sup>iii</sup> —Ca1—O2 <sup>i</sup>	80.9 (1)
O1—Ca1—O1 <sup>iii</sup>	81.6 (1)	O1 <sup>iii</sup> —Ca1—O1 <sub>w</sub>	95.9 (1)
O1—Ca1—O2	52.0 (1)	O1 <sup>ii</sup> —Ca1—O1 <sub>w</sub> <sup>i</sup>	155.5 (1)
O1—Ca1—O2 <sup>i</sup>	142.4 (1)	O2—Ca1—O2 <sup>i</sup>	144.7 (1)
O1—Ca1—O1 <sub>w</sub>	76.6 (1)	O2—Ca1—O1 <sub>w</sub>	77.0 (1)
O1—Ca1—O1 <sub>w</sub> <sup>i</sup>	130.1 (1)	O2—Ca1—O1 <sub>w</sub> <sup>i</sup>	78.2 (1)
O1 <sup>ii</sup> —Ca1—O1 <sup>iii</sup>	88.7 (1)	O1 <sub>w</sub> —Ca1—O1 <sub>w</sub> <sup>i</sup>	89.9 (1)

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



**Figure 2**  
 ORTEPII plot of the chain motif; the uncoordinated water molecules are not shown.

**Table 2**  
 Hydrogen-bond 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>
O5—H5 <sub>o</sub> ...O2 <sup>iv</sup>	0.85	1.89	2.720 (2)	166
O1 <sub>w</sub> —H1 <sub>w</sub> 1...O2 <sub>w</sub>	0.85	2.09	2.925 (8)	168
O1 <sub>w</sub> —H1 <sub>w</sub> 2...O4 <sup>v</sup>	0.85	1.91	2.704 (2)	156

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

The carbon-bound H atoms were positioned geometrically [C—H 0.93 or 0.97 Å] and were included in the refinement in the riding-model approximation, with  $U(H) = 1.2U_{eq}(C)$ . The H atoms of the coordinated water molecule were similarly treated [O—H 0.85 Å,  $U(H) = 1.5U_{eq}(O)$ ]; these were rotated to fit the electron density.

The uncoordinated water molecule is disordered over two positions and the occupancy refined to 0.67 (1):0.33 (1). The displacement parameters of the two components were set to equal each other; these were restrained to be nearly isotropic. H atoms were not added to the disordered components.

Data collection: *RAPID-AUTO* (Rigaku, 1998); cell refinement: *RAPID-AUTO*; data reduction: *CrystalStructure* (Rigaku/MS, 2002); program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *ORTEPII* (Johnson, 1976); software used to prepare material for publication: *SHELXL97*.

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