Received 24 May 2006

Accepted 21 June 2006

Acta Crystallographica Section E Structure Reports Online

ISSN 1600-5368

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Key indicators

Single-crystal X-ray study T = 120 K Mean σ (C–C) = 0.005 Å R factor = 0.038 wR factor = 0.095 Data-to-parameter ratio = 20.7

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

Poly[[tetraaquatris(monomethyl fumarato)distrontium(II)] monomethyl fumarate] at 120 K

The title compound, $\{[Sr_2(C_5H_5O_4)_3(H_2O)_4](C_5H_5O_4)\}_n$, crystallizes with three methyl fumarate ions and four water molecules coordinating the two independent strontium(II) ions. The coordination polyhedra are interconnected by edge-sharing to form chains, which are connected by hydrogen bonds into layers. The layers, in turn, are stacked with the fourth methyl fumarate anion intercalated.

Comment

Recent preclinical and clinical investigations have revealed that strontium(II) reduces bone resorption while at the same time it provides stimulation for new bone formation (Marie *et al.*, 2001). These investigations have led to a growing interest in strontium(II) salts, their crystal structures and synthetic methods that may provide products of high yield and purity (Christgau *et al.*, 2005; Stahl *et al.*, 2006). The present paper presents the structural investigation of such a new strontium(II) salt, (I).



The two independent strontium(II) ions are both eightcoordinated by three methyl fumarate ions and four water molecules (Table 1 and Fig. 1). The coordination polyhedra are interconnected by edge-sharing to form chains in the *a*axis direction (Fig. 2). Hydrogen bonds (Table 2) between water molecules and to the second, Sr-coordinated, methyl fumarate anion connect the chains into layers in the *ab* plane. Additional hydrogen bonds bind the uncoordinated methyl fumarate to these layers. The layers, in turn, are connected by electrostatic and van der Waals interactions between the layers including the fourth methyl fumarate anion. Three of the methyl fumarate ions are approximately planar, while in the fourth methyl fumarate ion the carboxylate group is twisted by approximately 60° (Table 3) to allow for its Sr coordination and packing.

Experimental

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The title compound was synthesized by sprinkling powdered strontium carbonate into a saturated solution of methyl fumarate in a

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molar ratio of 1:2 at 313 K. More methyl fumarate was added after cessation of the initial effervescence, followed by addition of strontium carbonate in a molar ratio of 1:2. The procedure was continued in this stepwise manner until a permanent precipitate was observed. Cooling to 293 K and filtration gave the title compound in high yield and purity.

Crystal data

$$\begin{split} & [\mathrm{Sr}_2(\mathrm{C}_5\mathrm{H}_5\mathrm{O}_4)_3(\mathrm{H}_2\mathrm{O})_4](\mathrm{C}_5\mathrm{H}_5\mathrm{O}_4) \\ & M_r = 763.66 \\ & \mathrm{Triclinic}, \ P\overline{1} \\ & a = 7.1336 \ (6) \\ & \dot{A} \\ & b = 10.7203 \ (8) \\ & \dot{A} \\ & c = 20.9384 \ (17) \\ & \dot{A} \\ & \alpha = 100.676 \ (2)^{\circ} \\ & \beta = 96.322 \ (2)^{\circ} \\ & \gamma = 107.096 \ (2)^{\circ} \end{split}$$

Data collection

Bruker SMART APEX diffractometer ω scans Absorption correction: multi-scan (*SADABS*; Sheldrick, 2002) $T_{\min} = 0.484, T_{\max} = 0.897$

Refinement

Refinement on F^2 $R[F^2 > 2\sigma(F^2)] = 0.038$ $wR(F^2) = 0.095$ S = 0.878421 reflections 407 parameters $V = 1480.7 (2) Å^{3}$ Z = 2 $D_{x} = 1.713 \text{ Mg m}^{-3}$ Mo K\alpha radiation $\mu = 3.69 \text{ mm}^{-1}$ T = 120 (2) KNeedle, colorless $0.23 \times 0.05 \times 0.03 \text{ mm}$

19750 measured reflections 8421 independent reflections 5753 reflections with $I > 2\sigma(I)$ $R_{\text{int}} = 0.057$ $\theta_{\text{max}} = 30.8^{\circ}$

H atoms treated by a mixture of independent and constrained refinement $w = 1/[\sigma^2(F_o^2) + (0.0363P)^2]$ where $P = (F_o^2 + 2F_c^2)/3$ $(\Delta/\sigma)_{max} = 0.001$ $\Delta\rho_{max} = 1.48 \text{ e } \text{Å}^{-3}$ $\Delta\rho_{min} = -0.71 \text{ e } \text{Å}^{-3}$

 Table 1

 Selected bond lengths (Å).

| Sr1-O11 | 2.499 (2) | Sr2-O4 | 2.569 (3) |
|-----------------------|-----------|------------------------|-----------|
| Sr1-O12 ⁱ | 2.509 (2) | Sr2-O32 ⁱⁱⁱ | 2.571 (2) |
| Sr1-O22 ⁱⁱ | 2.537 (2) | Sr2-O31 | 2.597 (2) |
| Sr1-O3 | 2.598 (3) | Sr2-O13 ^{iv} | 2.600 (2) |
| Sr1-O32 | 2.601 (2) | Sr2-O2 | 2.621 (2) |
| Sr1-O21 | 2.617 (2) | Sr2-O11 | 2.664 (2) |
| Sr1-O31 | 2.775 (2) | Sr2-O1 | 2.671 (2) |
| Sr1-O22 | 2.807 (2) | Sr2-O12 | 2.812 (2) |

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

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

| $D - H \cdot \cdot \cdot A$ | D-H | $H \cdot \cdot \cdot A$ | $D \cdot \cdot \cdot A$ | $D - \mathbf{H} \cdot \cdot \cdot A$ |
|---------------------------------------|----------------------|-------------------------|-------------------------|--------------------------------------|
| $O1-H1A\cdots O42^{v}$ | 0.834 (18) | 1.850 (19) | 2.681 (3) | 174 (4) |
| $O1 - H1B \cdot \cdot \cdot O21^{vi}$ | 0.818 (18) | 2.02 (2) | 2.813 (3) | 163 (4) |
| $O2-H2A\cdots O21^{vi}$ | 0.809 (18) | 2.017 (19) | 2.822 (3) | 174 (4) |
| $O2-H2B\cdots O41^{vi}$ | 0.837 (18) | 1.835 (19) | 2.667 (3) | 173 (4) |
| $O3-H3A\cdotsO1^{i}$ | 0.810 (18) | 1.98 (2) | 2.775 (4) | 169 (4) |
| $O3-H3B\cdots O2$ | 0.792 (18) | 2.03 (2) | 2.819 (4) | 171 (4) |
| $O4-H4A\cdots O41^{vi}$ | 0.821 (19) | 2.05 (3) | 2.788 (4) | 149 (4) |
| $O4-H4B\cdots O42^{v}$ | 0.807 (18) | 2.07 (2) | 2.835 (4) | 159 (4) |
| Symmetry codes: -x+1, -y+1, -z+1 | (i) $x - 1, y$ 1. | v, z; (v) – | -x+2, -y+1, - | -z + 1; (vi) |

023 012* 022 024 C23 013 C32 02 C14 C45 C13 03 O11 C12 C13 C33 013 031 034 012 042 032 Ô 01

Figure 1

The asymmetric unit of (I), showing 75% probability displacement ellipsoids and the atomic numbering. H atoms have been omitted for clarity. O12*, O22*, O13* and O32* are added to complete the Sr coordination, with symmetry codes (*cf* Table 1) i, ii, iv and iii, respectively.



Figure 2

The crystal packing of (I), viewed down the a axis. The eight-coordinate Sr atoms are shown as polyhedra. H atoms have been omitted for clarity.

Table 3

Selected methyl fumarate torsion angles (°).

| O11-C11-C12-C13 | -122.1 (4) | O31-C31-C32-C33 | -5.7 (5) |
|-----------------|------------|-----------------|------------|
| O13-C14-C13-C12 | -167.8(3) | O33-C34-C33-C32 | 7.1 (6) |
| C15-O14-C14-C13 | -179.7(3) | C35-O34-C34-C33 | -177.5(3) |
| O21-C21-C22-C23 | 10.8 (5) | O41-C41-C42-C43 | 12.5 (5) |
| 023-C24-C23-C22 | 10.8 (6) | O43-C44-C43-C42 | 9.2 (6) |
| C25-O24-C24-C23 | 177.6 (3) | C45-O44-C44-C43 | -178.8 (3) |
| | | | |

All H parameters were initially refined freely. In the final cycles the H atoms of the CH groups were placed in calculated positions with C-H = 0.93 Å and refined as riding atoms. The CH₃ groups were placed in calculated positions and the torsion angle allowed to refine

(C-H = 0.96 Å). For the water molecules, the O-H distances were restrained to 0.82 (2) Å. The displacement parameters were set to 1.2 (CH and CH₃) or 1.5 (OH) times U_{eq} of the corresponding C or O atoms. The highest peak is located 0.83 Å from atom Sr1.

Data collection: *SMART* (Bruker, 1999); cell refinement: *SAINT-Plus* (Bruker, 1999); data reduction: *SAINT-Plus* and *SADABS* (Sheldrick, 2002); program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *ORTEP-3* (Farrugia, 1997) and *ATOMS* (Dowty, 2000); software used to prepare material for publication: *SHELXL97*.

Data collection by Ms L. Berring and Ms A. Schøneberg are gratefully acknowledged.

References

- Bruker (1999). SMART and SAINT-Plus. Bruker AXS Inc., Madison, Wisconsin, USA.
- Christgau, S., Odderhede, J., Stahl, K. & Andersen, J. E. T. (2005). Acta Cryst. C61, m259–m262.
- Dowty, E. (2000). ATOMS. Version 5.1. Shape Software, Kingsport, Tennessee, USA.
- Farrugia, L. J. (1997). J. Appl. Cryst. 30, 565.
- Marie, P. J., Ammann, P., Boivin, G. & Rey, C. (2001). Calcif. Tissue Int. 69, 121-129.
- Sheldrick, G. M. (1997). SHELXS97 and SHELXL97. University of Göttingen, Germany.
- Sheldrick, G. M. (2002). SADABS. Version 2.03. University of Göttingen, Germany.
- Stahl, K., Andersen, J. E. T. & Christgau, S. (2006). Acta Cryst. C62, m144– m149.