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Key indicators

Single-crystal X-ray study T = 273 KMean $\sigma(C-C) = 0.001 \text{ Å}$ R factor = 0.024wR factor = 0.071 Data-to-parameter ratio = 32.1

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[2-(Dimethylsulfonio)-1-hydroxy-1-phosphonoethyl]phosphonate monohydrate

In the title crystal structure, C₄H₁₂O₇P₂S·H₂O, the sulfonium groups have a pyramidal geometry and bridging water molecules form a complex three-dimensional hydrogen-bond network involving neighboring phosphonate groups.

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Comment

The title compound, (I), belongs to a family of widely used compounds, bisphosphonates, with a characteristic P-C-P linkage that mimics the P-O-P linkage of inorganic diphosphate. Bisphosphonates are used to treat bone resorption diseases such as osteoporosis (Sambrook et al., 2004), and Paget's disease (Vasireddy et al., 2003). In addition, they have been found to have antiparasitic (Yardley et al., 2002; Martin et al., 2001), as well as anticancer activity (via $\gamma\delta$ T cells) (Sato et al., 2005). They act by targeting the mevalonate pathway enzyme farnesyl diphosphate synthase (FPPS) (EC 2.5.1.10) (Martin et al., 1999). Most bisphosphonates contain positively charged nitrogen-containing (ammonium, pyridinium, imidazolium) side chains, but other isosteres also have activity and we report here the structure of a novel sulfonium bisphosphonate, viz. (I).

$$\begin{array}{c} O \\ -OH \\ -S \\ OH \\ OH \\ HO \\ OH \\ (I) \end{array}$$

The sulfonium bisphosphonate crystallizes as a monohydrate and has one neutral and one monoanionic phosphonate group balancing the +1 charge on the sulfonium group. The PCP backbone of the bisphosphonate group exists in a conformation similar to those reported previously [P1-C1- $P2 = 113.87 (4)^{\circ}$ [incadronate (INC), isozoledronate (ISZ) and three hydrate forms of risedronate, namely the monohydrate (RMH), dihydrate (RDH) and 2.5-hydrate (RHP)] [INC 115.0 (2)°, ISZ 114.8 (1)°, RHP 112.4 (2)°, RDH 113.30 (15)° and RMH 113.22 (13)°; Montalvetti et al., 2003; Gossman et al., 2002, 2003]. The P-O distances are given in Table 1.

The dimethylsulfonium group has a distorted tetrahedral geometry with the two methyl groups having very similar C-S-C angles of $\sim 101^{\circ}$ [C4-S1-C3 = 101.04 (5)°, C4-S1- $C2 = 102.12 (5)^{\circ}$ and $C3 - S1 - C2 = 99.85 (4)^{\circ}$]. The distances of the two phosphate groups to the S atom $[S1 \cdots P1]$ = 3.6596 (3) Å and $S1 \cdot \cdot \cdot P2 = 4.3143$ (3) Å] are consistent with

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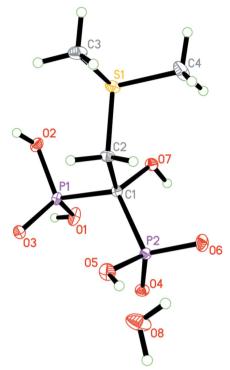


Figure 1SHELXTL (Bruker, 2001) plot showing 35% probability ellipsoids for non-H atoms and circles of arbitrary size for H atoms.

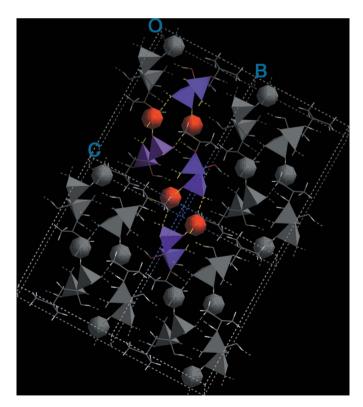


Figure 2CERIUS² (Accelrys, 2005) view of the crystal structure, showing the proposed hydrogen-bond interactions between neighboring molecules. Several such interactions occur by way of 'bridging' water molecules and have been highlighted to 'guide the eye'. Hydrogen bonds are represented by dashed yellow lines, water molecules as red spheres and phosphonate groups as purple polyhedra.

the electrostatic interaction between the positively charged S atom and the anionic phosphonate group. The water molecules form a complex hydrogen bond network with all adjacent phosphonate groups (Fig. 2).

Experimental

Bromoacetic acid (2 mmol) was added to a solution of dimethyl sulfide (2 mmol) in acetone (5 ml) and stirred for 2 h. The white precipitate was filtered off, washed with diethyl ether and dried in vacuo. The resulting precipitate was added to a mixture of H₃PO₃ (5 equivalents) and toluene (5 ml) and heated to 353 K until the mixture melted. POCl₃ (5 equivalents) was added slowly and the mixture stirred at 353 K for 4 h. Upon cooling, the supernatant was decanted and 4 ml water added. The mixture was refluxed for 1 h. Most of the solvent was then removed in vacuo and acetone was added to precipitate the anhydrous sulfonium bisphosphonate. The resulting white powder was collected and crystallized from ethanol-water (2:1). Analysis calculated for C₄H₁₂O₇P₂S: C 18.05, H 4.54%; found: C 18.43, H 4.70%. ¹H NMR (500 MHz, D₂O): δ 3.50–3.61 (*m*, 2H), 2.65 (s, 6H). ³¹P NMR (162 MHz, D₂O): δ 15.6 (s). The final crystals were grown by vapor diffusion of ethanol into an aqueous solution of the bisphosphonate at room temperature, using the sitting-drop method and yielding the monohydrate.

Crystal data

$C_4H_{12}O_7P_2S\cdot H_2O$	$D_x = 1.756 \text{ Mg m}^{-3}$
$M_r = 284.15$	Mo $K\alpha$ radiation
Monoclinic, $P2_1/c$	Cell parameters from 7336
a = 7.1107 (2) Å	reflections
b = 10.2149 (3) Å	$\theta = 2.9 - 36.3^{\circ}$
c = 15.1091 (4) Å	$\mu = 0.62 \text{ mm}^{-1}$
$\beta = 101.653 \ (10)^{\circ}$	T = 273 (2) K
$V = 1074.83 (6) \text{ Å}^3$	Column, colorless
Z = 4	$0.51 \times 0.24 \times 0.03 \text{ mm}$

Data collection

Bruker Kappa-APEXII CCD	5196 independent reflections
diffractometer	4605 reflections with $I > 2\sigma(I)$
φ and ω scans	$R_{\rm int} = 0.021$
Absorption correction: integration	$\theta_{\rm max} = 36.3^{\circ}$
(SHELXTL/XPREP;	$h = -11 \rightarrow 11$
Bruker, 2001)	$k = -15 \rightarrow 17$
$T_{\min} = 0.774, T_{\max} = 0.955$	$l = -13 \rightarrow 25$
26896 measured reflections	

Refinement

•	
Refinement on F^2	$w = 1/[\sigma^2(F_0^2) + (0.0343P)^2]$
$R[F^2 > 2\sigma(F^2)] = 0.024$	+ 0.3181P]
$wR(F^2) = 0.071$	where $P = (F_0^2 + 2F_c^2)/3$
S = 1.06	$(\Delta/\sigma)_{\text{max}} = 0.002$
5196 reflections	$\Delta \rho_{\text{max}} = 0.68 \text{ e Å}^{-3}$
162 parameters	$\Delta \rho_{\min} = -0.55 \text{ e Å}^{-3}$
H-atom parameters constrained	

Table 1 Selected bond lengths (Å).

P2-O6	1.5031 (7)	P1-O3	1.4962 (7)
P2-O4	1.5148 (7)	P1-O1	1.5335 (7)
P2-O5	1.5525 (7)	P1-O2	1.5624 (7)
P2-C1	1.8528 (8)	P1-C1	1.8486 (8)

organic papers

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

$D-H\cdots A$	D-H	$H \cdot \cdot \cdot A$	$D \cdot \cdot \cdot A$	D $ H$ $\cdot \cdot \cdot A$
$\begin{array}{c} \hline \\ O8-H13\cdots O3^{i} \\ O7-H4\cdots O3^{ii} \\ O2-H2\cdots O4^{iii} \\ O1-H1\cdots O6^{iv} \\ O5-H3\cdots O8 \\ O8-H14\cdots O4^{v} \\ \end{array}$	0.859 (14)	1.892 (15)	2.7435 (11)	171.0 (19)
	0.826 (12)	1.915 (12)	2.7138 (9)	162.6 (14)
	0.809 (13)	1.721 (13)	2.5287 (10)	176.4 (16)
	0.814 (13)	1.643 (13)	2.4420 (9)	166.4 (17)
	0.781 (13)	1.728 (13)	2.5049 (10)	172.7 (18)
	0.867 (14)	1.959 (15)	2.7975 (11)	162.3 (18)

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

Methyl H atom positions, $R-\mathrm{CH_3}$, were optimized by rotation about $R-\mathrm{C}$ bonds with idealized $C-\mathrm{H}$, $R-\mathrm{H}$ and $\mathrm{H}\cdots\mathrm{H}$ distances (methyl $C-\mathrm{H}=0.96$ Å with AFIX 137). Methylene and hydroxyl H-atom positions were located in late difference Fourier maps and restrained to ideal bond lengths ($O-\mathrm{H}=0.84$ Å) using an effective standard deviation of 0.02 Å. Methyl and hydroxyl H-atom $U_{\rm iso}(\mathrm{H})$ values were assigned as 1.5 times $U_{\rm eq}$ of the carrier atom; remaining H-atom $U_{\rm iso}(\mathrm{H})$ values were assigned as 1.2 times carrier atom $U_{\rm eq}$.

Data collection: *APEX2* (Bruker, 2004); cell refinement: *SAINT* (Bruker, 2001); data reduction: *SAINT*; program(s) used to solve structure: *SHELXTL* (Bruker, 2001); program(s) used to refine structure: *SHELXTL*; molecular graphics: *SHELXTL*; software used to prepare material for publication: XCIF (Bruker, 2001).

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