

[2-(3-Fluoropyridinium-1-yl)-1-hydroxy-1-phosphonoethyl]phosphonate**Rong Cao,^a Michael P. Hudock,^a
Yonghui Zhang,^b Scott R.
Wilson^c and Eric Oldfield^{b*}**^aCenter for Biophysics and Computational Biology, University of Illinois at Urbana-Champaign, 607 South Mathews Avenue, Urbana, Illinois 61801, USA, ^bDepartment of Chemistry, University of Illinois at Urbana-Champaign, 600 South Mathews Avenue, Urbana, Illinois 61801, USA, and ^cSchool of Chemical Sciences, Box 59-1, University of Illinois at Urbana-Champaign, 505 South Mathews Avenue, Urbana, Illinois 61801, USA

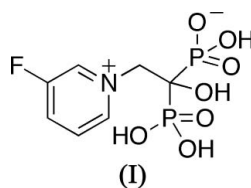
Correspondence e-mail: eo@chad.scs.uiuc.edu

Key indicatorsSingle-crystal X-ray study
 $T = 193$ K
Mean $\sigma(C-C) = 0.003$ Å
 R factor = 0.032
 wR factor = 0.088
Data-to-parameter ratio = 18.2For details of how these key indicators were automatically derived from the article, see <http://journals.iucr.org/e>.

The title compound, $C_7H_{10}FNO_7P_2$, crystallizes in the zwitterionic form. In the crystal structure, molecules are linked *via* intermolecular $O-H \cdots O$ hydrogen bonds involving phosphonate groups, forming a two-dimensional framework. In addition, weak intermolecular $C-H \cdots O$ hydrogen bonds involving the pyridinium groups further connect molecules, forming a three-dimensional framework.

Comment

Nitrogen-containing geminal bisphosphonates are used clinically to treat a variety of bone resorption diseases such as osteoporosis (Sambrook *et al.*, 2004) and Paget's disease (Vasireddy *et al.*, 2003) and more recently they have been found to have antiparasitic (Yardley *et al.*, 2002), anticancer (Cleardin, 2005) and immunomodulatory activity (Sanders *et al.*, 2004). It is believed that they act by inhibiting the isoprenoid biosynthesis pathway enzyme farnesyl diphosphate synthase (FPPS) (EC 2.5.1.10) (Martin *et al.*, 1999; Hosfield *et al.*, 2004; van Beek *et al.*, 1999) with the charged (ammonium, imidazolium) side chains acting as carbocation transition state reactive intermediates. Another class of bisphosphonates are the pyridinium-1-yl species, which contain a fixed (+1) side-chain charge (Sanders *et al.*, 2005). We report here the first structure of one such active compound, *viz.* (I).



The molecule of (I) crystallizes in the zwitterionic form and there are no solvent molecules present (unlike the monohydrates found with the sulfonium, phosphonium and arsonium bisphosphonates; Zhang *et al.*, 2006; Cao *et al.*, 2006; Hudock *et al.*, 2006). Consequently, there must be three protonated phosphonate O atoms (found to be O1, O4 and O5) and three non-protonated phosphonate O atoms (found to be O2, O3 and O6), as shown in Fig. 1.

The $P1 \cdots N1$ distance [$3.2469(13)$ Å] is considerably shorter than the $P2 \cdots N1$ distance [$4.1195(14)$ Å], consistent with a strong intramolecular interaction between the pyridinium N atom and the anionic phosphonate group, and the overall structure closely resembles that found for the monohydrate form of risedronate (Gossman *et al.*, 2003) which contains a pyridinium N atom (at a position equivalent to that found here for C4). In the crystal structure, molecules are

Received 17 January 2006

Accepted 6 February 2006

linked *via* intermolecular O—H···O hydrogen bonds involving phosphonate groups, forming a two-dimensional framework (Fig. 2 and Table 1). In addition, weak intermolecular C—H···O hydrogen bonds involving the pyridinium groups further connect molecules, forming a three-dimensional framework (Table 1).

Experimental

The title compound was prepared as described previously (Sanders *et al.*, 2005). Crystals were grown by vapor diffusion of ethanol into an aqueous solution of the bisphosphonate at room temperature using the sitting-drop method.

Crystal data

$C_7H_{10}FNO_7P_2$	$D_x = 1.810 \text{ Mg m}^{-3}$
$M_r = 301.10$	Mo $K\alpha$ radiation
Monoclinic, $P2_1/c$	Cell parameters from 4398 reflections
$a = 6.1892 (10) \text{ \AA}$	$\theta = 3.1\text{--}29.9^\circ$
$b = 19.1693 (4) \text{ \AA}$	$\mu = 0.44 \text{ mm}^{-1}$
$c = 9.3312 (10) \text{ \AA}$	$T = 193 (2) \text{ K}$
$\beta = 93.667 (10)^\circ$	Block (acircular), colourless
$V = 1104.8 (3) \text{ \AA}^3$	$0.10 \times 0.10 \times 0.06 \text{ mm}$
$Z = 4$	

Data collection

Bruker Kappa-APEXII CCD diffractometer	3201 independent reflections
φ and ω scans	2665 reflections with $I > 2\sigma(I)$
Absorption correction: integration (SHELXTL/XPREP; Bruker, 2001)	$R_{\text{int}} = 0.032$
$T_{\text{min}} = 0.967$, $T_{\text{max}} = 0.981$	$\theta_{\text{max}} = 30.0^\circ$
15680 measured reflections	$h = -8 \rightarrow 8$
	$k = -26 \rightarrow 26$
	$l = -13 \rightarrow 13$

Refinement

Refinement on F^2	$w = 1/[\sigma^2(F_o^2) + (0.0429P)^2 + 0.6018P]$
$R[F^2 > 2\sigma(F^2)] = 0.033$	where $P = (F_o^2 + 2F_c^2)/3$
$wR(F^2) = 0.088$	$(\Delta/\sigma)_{\text{max}} = 0.002$
$S = 1.03$	$\Delta\rho_{\text{max}} = 0.50 \text{ e \AA}^{-3}$
3201 reflections	$\Delta\rho_{\text{min}} = -0.32 \text{ e \AA}^{-3}$
176 parameters	
H atoms treated by a mixture of independent and constrained refinement	

Table 1

Hydrogen-bond geometry (\AA , $^\circ$).

$D\text{---}H\cdots A$	$D\text{---}H$	$H\cdots A$	$D\cdots A$	$D\text{---}H\cdots A$
O5—H3···O3 ⁱ	0.867 (15)	1.577 (16)	2.4209 (15)	163 (2)
O1—H1···O6 ⁱⁱ	0.834 (15)	1.679 (16)	2.4950 (15)	165 (2)
O4—H2···O2 ⁱⁱⁱ	0.840 (15)	1.749 (15)	2.5846 (15)	173 (2)
O7—H4···O2 ⁱⁱ	0.827 (15)	1.880 (15)	2.7056 (15)	177 (2)
C2—H5···O5 ⁱ	0.99	2.40	3.3326 (17)	156
C2—H6···O3 ⁱⁱⁱ	0.99	2.26	3.2216 (16)	163
C3—H7···O1 ⁱⁱⁱ	0.95	2.51	3.430 (2)	164
C5—H8···O4 ^{iv}	0.95	2.59	3.312 (2)	133
C6—H9···O6 ^v	0.95	2.45	3.260 (2)	143

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

Methyl H-atom positions, $R\text{---}CH_3$, were optimized by rotation about $R\text{---}C$ bonds with idealized $C\text{---}H$, $R\text{---}H$ and $H\cdots H$ distances (methyl $C\text{---}H = 0.96 \text{ \AA}$ with AFIX). Hydroxyl H-atom positions were located in late difference Fourier maps and restrained to ideal bond

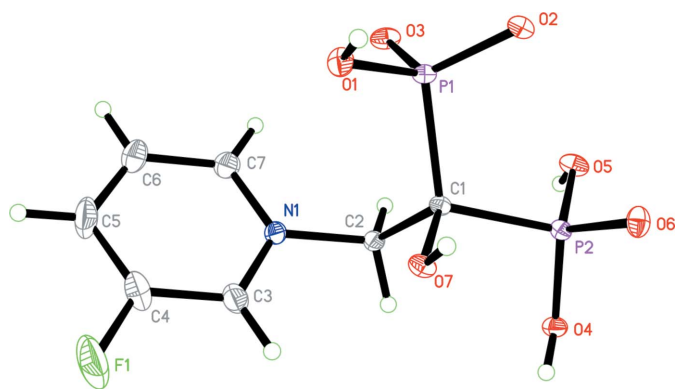


Figure 1

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

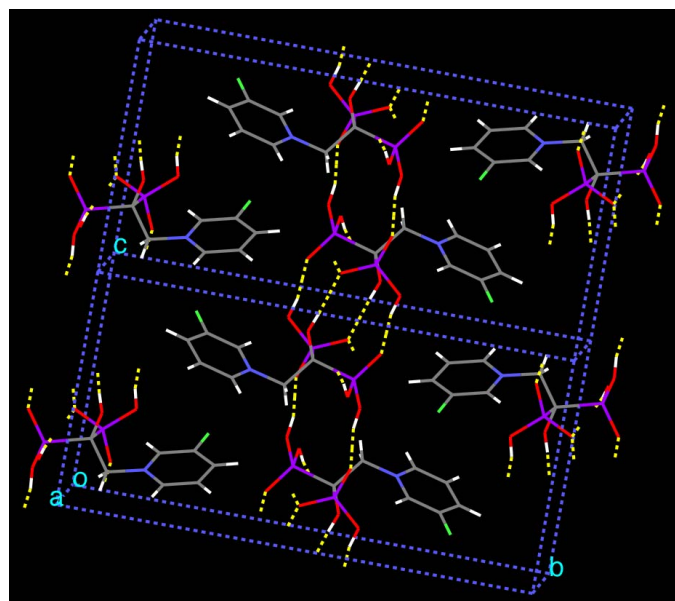


Figure 2

CERIU2 (Accelrys, 2005) view of the crystal structure, showing the hydrogen-bonded network between the bisphosphonates. Hydrogen bonds are shown as dashed yellow lines.

lengths ($O\text{---}H = 0.84 \text{ \AA}$) using an effective standard deviation of 0.02 \AA . The remaining H atoms were included as idealized riding atoms (methylene $C\text{---}H = 0.97 \text{ \AA}$ and ring $C\text{---}H = 0.93 \text{ \AA}$). Methyl and hydroxyl H-atom $U_{\text{iso}}(H)$ values were assigned as 1.5 times U_{eq} of the carrier atom; remaining H-atom $U_{\text{iso}}(H)$ values were assigned as 1.2 times carrier U_{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).

This work was supported in part by the United States Public Health Service (grant No. GM-073216 to EO). YZ is an American Heart Association, Midwest Affiliate, Postdoctoral Fellow. The Materials Chemistry Laboratory at the University of Illinois was supported in part by grants NSF CHE 95-03145

and NSF CHE 03–43032 from the National Science Foundation.

References

- Accelrys (2005). *CERIUS²*. Accelrys Inc., San Diego, CA, USA.
- Beek, E. van, Pieterman, E., Cohen, L., Lowik, C. & Papapoulos, S. (1999). *Biochem. Biophys. Res. Commun.* **255**, 491–494.
- Bruker (2001). *SAINT* (Version 6.22), *SHELXTL* (Version 6.12) and *XCIF*. Bruker AXS Inc., Madison, Wisconsin, USA.
- Bruker (2004). *APEX2*. Version 1.0-27. Bruker AXS Inc., Madison, Wisconsin, USA.
- Cao, R., Hudock, M. P., Zhang, Y., Wilson, S. R. & Oldfield, E. (2006). *Acta Cryst.* **E62**. Submitted.
- Clezardin, P. (2005). *Cancer Treat. Rev.* **31S3**, 1–8.
- Gossman, W. L., Wilson, S. R. & Oldfield, E. (2003). *Acta Cryst.* **C59**, m33–m36.
- Hosfield, D. J., Zhang, Y., Dougan, D. R., Broun, A., Tari, L. W., Swanson, R. V. & Finn, J. (2004). *J. Biol. Chem.* **279**, 8526–8529.
- Hudock, M. P., Cao, R., Zhang, Y., Wilson, S. R. & Oldfield, E. (2006). *Acta Cryst.* **E62**, o843–o845.
- Martin, M. B., Arnold, W., Heath, H. T. III, Urbina, J. A. & Oldfield, E. (1999). *Biochem. Biophys. Res. Commun.* **263**, 754–758.
- Sambrook, P. N., Geusens, P., Ribot, C., Solimano, J. A., Ferrer-Barriendos, J., Gaines, K., Verbruggen, N. & Melton, M. E. (2004). *J. Intern. Med.* **255**, 503–511.
- Sanders, J. M., Ghosh, S., Chan, J. M. W., Meints, G., Wang, H., Raker, A. M., Song, Y. C., Colantino, A., Burzynska, A., Kafarski, P., Morita, C. T. & Oldfield, E. (2004). *J. Med. Chem.* **47**, 375–384.
- Sanders, J. M., Song, Y., Chan, J. M., Zhang, Y., Jennings, S., Kosztowski, T., Odeh, S., Flessner, R., Schwerdtfeger, C., Kotsikorou, E., Meints, G. A., Gomez, A. O., Gonzalez-Pacanowska, D., Raker, A. M., Wang, H., van Beek, E. R., Papapoulos, S. E., Morita, C. T. & Oldfield, E. (2005). *J. Med. Chem.* **48**, 2957–2963.
- Vasireddy, S., Talwalkar, A., Miller, H., Mehan, R. & Swinson, D. R. (2003). *Clin. Rheumatol.* **22**, 376–380.
- Yardley, V., Khan, A. A., Martin, M. B., Slifer, T. R., Araujo, F. G., Moreno, S. N., Docampo, R., Croft, S. L. & Oldfield, E. (2002). *Antimicrob. Agents Chemother.* **46**, 929–931.
- Zhang, Y., Cao, R., Hudock, M. P., Wilson, S. R. & Oldfield, E. (2006). *Acta Cryst.* **E62**, o1006–o1008.