

C(32)	-0.0311 (9)	-0.1137 (10)	0.3969 (5)	0.027 (2)
C(33)	0.0095 (10)	-0.1733 (9)	0.4677 (6)	0.030 (2)
C(34)	0.1407 (10)	-0.1622 (10)	0.4977 (6)	0.032 (2)
C(35)	0.2327 (10)	-0.0877 (11)	0.4566 (6)	0.037 (3)
C(36)	0.1955 (9)	-0.0240 (10)	0.3873 (6)	0.030 (2)

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Table 2. Selected geometric parameters (Å, °)

Pb(1)—C(31)	2.201 (9)	C(21)—C(22)	1.363 (13)
Pb(1)—C(21)	2.213 (8)	C(21)—C(26)	1.392 (13)
Pb(1)—C(1)	2.214 (8)	C(22)—C(23)	1.397 (13)
Pb(1)—Br(1)	2.985 (2)	C(23)—C(24)	1.39 (2)
Pb(1)—Br(1a)	2.885 (2)	C(24)—C(25)	1.36 (2)
C(1)—C(11)	1.484 (12)	C(25)—C(26)	1.387 (14)
C(11)—C(16)	1.396 (14)	C(31)—C(36)	1.375 (12)
C(11)—C(12)	1.406 (14)	C(31)—C(32)	1.388 (12)
C(12)—C(13)	1.370 (14)	C(32)—C(33)	1.383 (13)
C(13)—C(14)	1.370 (15)	C(33)—C(34)	1.361 (14)
C(14)—C(15)	1.38 (2)	C(34)—C(35)	1.383 (14)
C(15)—C(16)	1.385 (13)	C(35)—C(36)	1.372 (14)
C(31)—Pb(1)—C(21)	115.4 (3)	Br(1)—Pb(1)—Br(1a <sup>i</sup> )	173.610 (15)
C(31)—Pb(1)—C(1)	127.8 (3)	Pb(1)—Br(1)—Pb(1b <sup>ii</sup> )	122.73 (5)
C(21)—Pb(1)—C(1)	116.7 (3)	C(11)—C(1)—Pb(1)	113.9 (6)
C(31)—Pb(1)—Br(1a <sup>i</sup> )	87.3 (2)	C(16)—C(11)—C(1)	120.1 (9)
C(21)—Pb(1)—Br(1a <sup>i</sup> )	93.6 (3)	C(12)—C(11)—C(1)	122.1 (9)
C(1)—Pb(1)—Br(1a <sup>i</sup> )	93.4 (3)	C(22)—C(21)—Pb(1)	121.6 (7)
C(31)—Pb(1)—Br(1)	88.6 (2)	C(26)—C(21)—Pb(1)	119.5 (7)
C(21)—Pb(1)—Br(1)	92.6 (3)	C(36)—C(31)—Pb(1)	116.9 (6)
C(1)—Pb(1)—Br(1)	85.3 (3)	C(32)—C(31)—Pb(1)	122.1 (6)

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

Systematic absences ( $h0l$ )  $h + l = 2n + 1$ , ( $00l$ )  $l = 2n + 1$  and ( $0k0$ )  $k = 2n + 1$  were detected. The structure was solved by standard Patterson and difference Fourier methods (*SHELXTL-Plus*; Sheldrick, 1987) and refined satisfactorily within space group  $P2_1/c$  (No. 14) by full-matrix least-squares calculations (*SHELXL*; Sheldrick, 1994). The H atoms were placed in geometrically calculated positions and refined with common isotropic displacement parameters for different C—H types (H<sub>aryl</sub>, H<sub>alkyl</sub>). Computer programs used were *SHELXTL-Plus* (Sheldrick, 1987), *SHELXL*, *PARST* (Nardelli, 1983), *PLATON* (Spek, 1990) and *MISSYM* (Le Page, 1987).

Financial support from Fonds der Chemischen Industrie is gratefully acknowledged.

Lists of structure factors, anisotropic displacement parameters, H-atom coordinates and least-squares-planes data have been deposited with the IUCr (Reference: SH1079). Copies may be obtained through The Managing Editor, International Union of Crystallography, 5 Abbey Square, Chester CH1 2HU, England.

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## A Hydrate of the Sodium Salt of a Penem Carboxylic Acid

YOSHIKI OYAMA, SEIICHI IMAJO, RIE TANAKA  
AND MASAJI ISHIGURO\*

*Suntory Institute for Biomedical Research, Shimamoto,  
Mishima, Osaka 618, Japan*

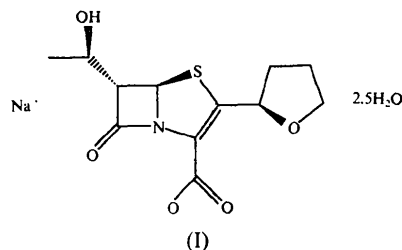
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## Abstract

The crystal structure of sodium (5*R*,6*S*)-6-[(*R*)-1-hydroxyethyl]-7-oxo-3-[(*R*)-2-tetrahydrofuryl]-4-thia-1-azabicyclo[3.2.0]hept-2-ene-2-carboxylate 2.5-hydrate, Na<sup>+</sup>.C<sub>12</sub>H<sub>14</sub>NO<sub>5</sub>S<sup>-</sup>. $\frac{5}{2}$ H<sub>2</sub>O, a member of a new class of  $\beta$ -lactam antibiotics, has been determined by X-ray diffraction. The C(3)—S(1) bond (bridgehead C atom) is longer than the C(2)—S(1) bond, although the latter is usually longer in penicillins and cephalosporins. The crystal packing shows a three-layer sheet in which two layers of the molecules are held together by Na<sup>+</sup> ions and by water molecules through a network of hydrogen bonds.

## Comment

The title compound (I), which has the tetrahydrofuran ring at the C(2) position, shows potent *in vivo* activity by oral administration (Nishino *et al.*, 1991). Knowledge of the accurate crystal structure of this compound should be useful for understanding the interaction of the molecule with water and biological macromolecules.



The bond lengths and angles of the penem nucleus and the carboxyl group of this molecule are similar to those of other penems. Those of the tetrahydrofuran ring are also similar to those found for tetrahydro-2-furancarboxylic acid (Bando, Takano, Tanaka, Iwata & Ishiguro, 1990). The  $\beta$ -lactam ring of the penem has a puckered structure like those of penicillins and cephalosporins. This is different from the conformations of unfused  $\beta$ -lactams, the rings of which are usually flat (Bando *et al.*, 1989). The angle C(3)—S(1)—C(2) of  $90.7(2)^\circ$  is smaller than the corresponding average angle of  $94.2(18)^\circ$  for 14 cephalosporins and is close to the average angle of  $90.6(8)^\circ$  for seven penicillins of the ampicillin type obtained through a search of the Cambridge Structural Database (Allen *et al.*, 1979). In the penem the C(3)—S(1) bond [ $1.831(4) \text{ \AA}$ ] is longer than the C(2)—S(1) bond [ $1.775(5) \text{ \AA}$ ], whereas in the penicillins and the cephalosporins the C(2)—S(1) bond is longer [averages  $1.85(1)$  and  $1.83(2) \text{ \AA}$ , respectively] than the C(3)—S(1) bond [average  $1.80(1) \text{ \AA}$  for each]. This characteristic difference in the bond lengths could

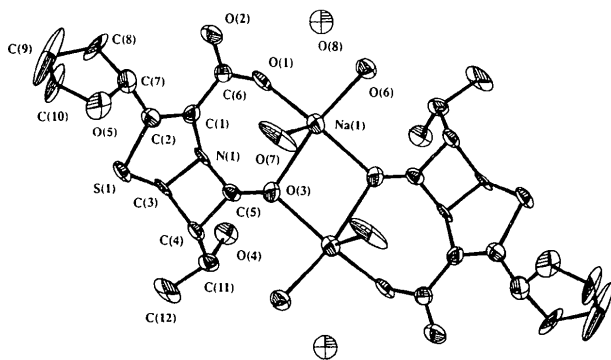


Fig. 1. ORTEP (Johnson, 1965) drawing of the molecular structure with atom-numbering scheme. The ellipsoids correspond to 50% probability levels of atomic displacement.

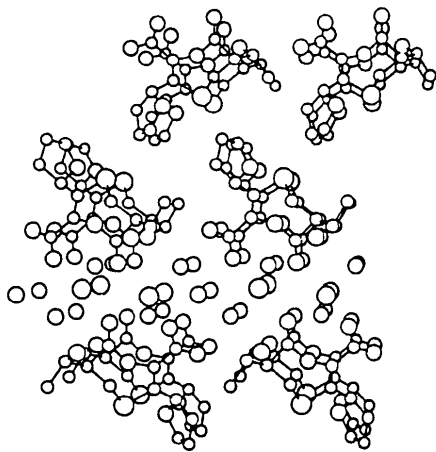


Fig. 2. PLUTO (Motherwell & Clegg, 1978) drawing of the crystal packing.

be the reason for the unusual reactivity of the penem in photolysis (Iwata, Tanaka, Oyama & Ishiguro, 1991), alkaline hydrolysis (Iwata, Tanaka & Ishiguro, 1990) and biological hydrolysis.

The C—O bonds of the carboxylate group are intermediate between single- and double-bonds. This indicates that the group is deprotonated and that electron delocalization occurs, although the C(6)—O(1) bond [ $1.281(6) \text{ \AA}$ ] is longer than the C(6)—O(2) bond by  $0.05 \text{ \AA}$ . The C(5)—O(3) bond [ $1.215(5) \text{ \AA}$ ] is slightly longer than the corresponding average bond length [ $1.19(2) \text{ \AA}$ ] for five penems obtained from the Cambridge Structural Database as a result of the coordination of O(1) and O(3) to Na(1).

The structure contains pairs of Na<sup>+</sup> cations, each pair being bridged by two  $\beta$ -lactam carbonyl groups. The distorted trigonal-bipyramidal coordination geometry around each Na<sup>+</sup> is completed by coordination to one of the carboxylate O atoms [O(1)] and to the O atoms [O(6) and O(7)] from two of the water molecules. These water molecules are also hydrogen bonded to the carbonyl, carboxylate and hydroxyethyl groups of the penem. The water O atom O(8) is hydrogen bonded to a carboxylate group and to O(6). The coordination to the Na<sup>+</sup> cations and the hydrogen bonds link two layers of penem molecules to form a three-layer sheet. There are no interactions between the sheets except for van der Waals contacts.

## Experimental

### Crystal data

Na<sup>+</sup>.C<sub>12</sub>H<sub>14</sub>NO<sub>5</sub>S<sup>-</sup>. $\frac{5}{2}$ H<sub>2</sub>O

$M_r = 352.3$

Orthorhombic

$P2_12_12$

$a = 9.208(1) \text{ \AA}$

$b = 32.454(2) \text{ \AA}$

$c = 5.495(3) \text{ \AA}$

$V = 1642.1(10) \text{ \AA}^3$

$Z = 4$

$D_x = 1.43 \text{ Mg m}^{-3}$

Cu K $\alpha$  radiation

$\lambda = 1.5418 \text{ \AA}$

Cell parameters from 45 reflections

$\theta = 19\text{--}32^\circ$

$\mu = 2.307 \text{ mm}^{-1}$

$T = 293 \text{ K}$

Hexagonal plate

$0.5 \times 0.3 \times 0.3 \text{ mm}$

Colourless

### Data collection

Rigaku AFC-5 diffractometer

$\omega/2\theta$  scans

Absorption correction:

none

1606 measured reflections

1606 independent reflections

1592 observed reflections

$[F > 3\sigma(F)]$

$\theta_{\max} = 63^\circ$

$h = 0 \rightarrow 11$

$k = 0 \rightarrow 38$

$l = 0 \rightarrow 6$

3 standard reflections

monitored every 100

reflections

intensity variation: 0.8%

### Refinement

Refinement on  $F$

$R = 0.059$

$wR = 0.110$

$(\Delta/\sigma)_{\max} = 4.84$

$\Delta\rho_{\max} = 0.51 \text{ e \AA}^{-3}$

$\Delta\rho_{\min} = -0.27 \text{ e \AA}^{-3}$

S = 1.92

1592 reflections

204 parameters

H atoms not located

$$w = 1/[\sigma^2(F_o) + 0.02081F_o + 0.0049F_o^2]$$

Atomic scattering factors  
from *International Tables*  
for *X-ray Crystallography*  
(1974, Vol. IV)

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Table 1. Fractional atomic coordinates and equivalent isotropic displacement parameters (Å<sup>2</sup>)

$$U_{eq} = (1/3)\sum_i\sum_j U_{ij}a_i^*a_j^*a_i \cdot a_j$$

	x	y	z	U <sub>eq</sub>
S(1)	0.0444 (1)	0.3203 (1)	0.8346 (2)	0.040
Na(1)	0.1888 (2)	0.4939 (1)	0.8123 (4)	0.039
O(1)	0.3208 (3)	0.4459 (1)	1.0278 (6)	0.037
O(2)	0.4625 (3)	0.4015 (1)	0.8265 (9)	0.051
O(3)	-0.0239 (3)	0.4535 (1)	0.9042 (6)	0.035
O(4)	-0.1491 (4)	0.4125 (1)	1.4301 (7)	0.044
O(5)	0.2486 (5)	0.2999 (1)	0.4651 (8)	0.058
O(6)	0.3569 (3)	0.5397 (1)	0.6595 (7)	0.040
O(7)	0.1504 (6)	0.4599 (2)	0.4524 (9)	0.086
O(8)	1/2	1/2	0.2681 (9)	0.047
N(1)	0.0889 (4)	0.3919 (1)	1.0352 (7)	0.030
C(1)	0.2221 (4)	0.3828 (1)	0.9117 (9)	0.032
C(2)	0.2119 (5)	0.3464 (1)	0.7922 (9)	0.037
C(3)	-0.0071 (4)	0.3566 (1)	1.0747 (8)	0.030
C(4)	-0.1405 (4)	0.3854 (1)	1.0266 (8)	0.030
C(5)	-0.0240 (5)	0.4179 (1)	0.9725 (7)	0.029
C(6)	0.3459 (5)	0.4116 (1)	0.9209 (9)	0.032
C(7)	0.3233 (6)	0.3255 (1)	0.639 (1)	0.045
C(8)	0.4265 (7)	0.2979 (2)	0.777 (2)	0.072
C(9)	0.398 (2)	0.2559 (3)	0.671 (3)	0.173
C(10)	0.317 (1)	0.2611 (2)	0.462 (2)	0.083
C(11)	-0.2373 (4)	0.3964 (1)	1.2395 (8)	0.034
C(12)	-0.3233 (6)	0.3592 (2)	1.327 (1)	0.062

Table 2. Selected geometric parameters (Å, °)

S(1)—C(2)	1.775 (5)	Na(1)···O(7)	2.292 (6)
S(1)—C(3)	1.831 (4)	O(6 <sup>ii</sup> )···O(2)	2.691 (6)
O(1)—C(6)	1.281 (6)	O(6 <sup>ii</sup> )···O(4 <sup>iii</sup> )	2.766 (5)
O(2)—C(6)	1.235 (7)	O(7)···O(1 <sup>iv</sup> )	2.848 (7)
O(3)—C(5)	1.215 (5)	O(7)···O(3)	2.963 (7)
Na(1)···O(1)	2.304 (4)	O(7)···O(4 <sup>iv</sup> )	3.160 (7)
Na(1)···O(3)	2.410 (4)	O(8)···O(1 <sup>iv</sup> )	2.748 (6)
Na(1)···O(3 <sup>i</sup> )	2.340 (4)	O(8)···O(6)	2.833 (6)
Na(1)···O(6)	2.305 (4)		
C(2)—S(1)—C(3)	90.7 (2)		

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

The high values of  $wR$  and  $(\Delta/\sigma)_{\max}$  result from disorder of the tetrahydrofuryl group, the hydroxyethyl group and the water O atom (O7). Data collection: *AFC-5 Data Collection Software* (Rigaku Corporation, 1984a). Data reduction: *RASA-5P* (Rigaku Corporation, 1984b). Program(s) used to solve structure: *MULTAN11/84* (Main, Germain & Woolfson, 1984). Program(s) used to refine structure: *RASA-5P*. Molecular graphics: *ORTEP* (Johnson, 1965); *PLUTO* (Motherwell & Clegg, 1978).

Lists of structure factors, anisotropic displacement parameters and complete geometry have been deposited with the IUCr (Reference: AS1108). Copies may be obtained through The Managing Editor, International Union of Crystallography, 5 Abbey Square, Chester CH1 2HU, England.

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## Di- $\mu$ -chloro-bis[dichlorotris(2-propanol)cerium(III)] at 153 K

MARTINA SCHÄFER AND REGINE HERBST-IRMER

*Institut für Anorganische Chemie, Universität Göttingen, Tammannstr. 4, 37077 Göttingen, Germany*

ULRICH GROTH AND THOMAS KÖHLER

*Institut für Organische Chemie, Universität Göttingen, Tammannstr. 2, 37077 Göttingen, Germany*

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### Abstract

The structure of di- $\mu$ -chloro-bis[dichlorotris(2-propanol)cerium(III)], [CeCl<sub>3</sub>(C<sub>3</sub>H<sub>8</sub>O)<sub>3</sub>]<sub>2</sub>, is reported. The Ce atoms are bridged by two Cl atoms and so a dimer is formed.

### Comment

The addition of trimethylsilylpropargyl† magnesium bromide to aldehyde affords, predominantly, allenyl alcohol. The regioselectivity changes dramatically when

† Propargyl = 2-propynyl.