

The results of this refinement are: number of refined parameters $p = 315$, number of reflections used in the LS refinement $n = 4035$, $wR_2 = 0.2711$ for 4035 reflections, $S = [\sum w(F_o^2 - F_c^2)/(p - n)]^{1/2} = 0.746$, $w = 1/[\sigma^2(F_o^2) + (0.1658 P)^2]$ where $P = 1/3[\max.(F_o^2, 0) + 2F_c^2]$, $wR_2' = 0.2796$ for all 4062 reflections (27 reflections with $\Delta/\sigma > 4$ omitted in the refinement), $R_1 = 0.0924$ for 1267 $F_o > 4\sigma(F_o)$, $R_1' = 0.2011$ for all data, $S = [\sum w(F_o - F_c)^2/(p - n)]^{1/2} = 0.827$ where $wR_2 = [\sum w(F_o^2 - F_c^2)^2/\sum w(F_o^2)^2]^{1/2}$, $R_1 = \sum |F_o - F_c|/\sum F_o$. A comparison of the structural parameters (coordinates, displacement parameters, bond distances, angles, etc.) from the two refinements shows that, as a result of the increased number of observations, the e.s.d.'s are much lower ($\sim 1/3$) for the F_o^2 refinement but that differences between bond distances and angles are not significant, which supports the view that the quality of intensity data is much more important than the amount of data. Half-normal probability plots and a table of the distances and angles obtained from the F_o^2 refinement have been deposited.

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Lists of structure factors, anisotropic thermal parameters, H-atom coordinates and complete geometry, together with principal axes of thermal ellipsoids, half-normal probability plots and distances and angles from the F_o^2 refinement, have been deposited with the British Library Document Supply Centre as Supplementary Publication No. SUP 55192 (20 pp.). Copies may be obtained through The Technical Editor, International Union of Crystallography, 5 Abbey Square, Chester CH1 2HU, England. [CIF reference: KA1012]

References

- Amlaiky, N., Leclerc, J., Decker, N. & Schwartz, J. (1983). *Eur. J. Med. Chem. Chim. Theor.* **18**, 437-439.
- Belletti, D., Uguzzoli, F., Cantoni, A. & Pasquinelli, G. (1979). *Gestione on Line di Diffratometro a Cristallo Singolo Siemens-AED con Sistema General Automation Jumbo 220*. Internal Report 1-3/79. Centro di Studio per la Strutturistica Diffratometrica del CNR, Parma, Italy.
- Cremer, D. & Pople, J. A. (1975). *J. Am. Chem. Soc.* **97**, 1354-1358.
- Jamart-Grégoire, B., Caubère, P., Blanc, M., Gnassounou, J. P. & Advenier, C. (1989). *J. Med. Chem.* **32**, 315-320.
- Johnson, C. K. (1965). *ORTEP*. Report ORNL-3794. Oak Ridge National Laboratory, Tennessee, USA.
- Lawton, S. L. & Jacobson, R. A. (1965). *The Reduced Cell and its Crystallographic Applications*. Ames Laboratory. Available from the Clearinghouse for Federal Scientific and Technical Information, National Bureau of Standards, US Department of Commerce, Springfield, Virginia, USA.
- Lehmann, M. S. & Larsen, F. K. (1974). *Acta Cryst.* **A30**, 580-589.
- Le Page, Y. (1987). *J. Appl. Cryst.* **A20**, 264-269.
- Main, P., Fiske, S. J., Hull, S. E., Lessinger, L., Germain, G., Declercq, J.-P. & Woolfson, M. M. (1980). *MULTAN80. A System of Computer Programs for the Automatic Solution of Crystal Structures from X-ray Diffraction Data*. Univs. of York, England, and Louvain, Belgium.
- Motherwell, W. D. S. & Clegg, W. (1978). *PLUTO*. Program for plotting molecular and crystal structures. Univ. of Cambridge, England.
- Mugnoli, A. (1985). *J. Appl. Cryst.* **18**, 183-184.
- Mugnoli, A. (1990). *CAVITY*. Program to locate and measure spherical voids in a crystal structure. Univ. of Genoa, Italy.
- Nardelli, M. (1983a). *Comput. Chem.* **7**, 95-98.

- Nardelli, M. (1983b). *Acta Cryst.* **C39**, 1141-1142.
- Nardelli, M. & Mangia, A. (1984). *Ann. Chim. (Rome)*, **74**, 163-174.
- Sheldrick, G. M. (1976). *SHELX76*. Program for crystal structure determination. Univ. of Cambridge, England.
- Spek, A. L. (1988). *J. Appl. Cryst.* **21**, 578-579.
- Zachariasen, W. H. (1963). *Acta Cryst.* **16**, 1139-1144.

Acta Cryst. (1992). **C48**, 1737-1739

Structure of a Novel Carbapenem Antibiotic, Meropenem

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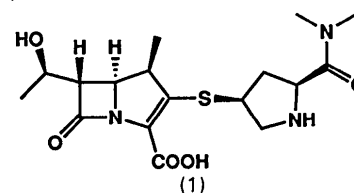
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Abstract

The absolute configuration of the title compound, (4*R*,5*S*,6*S*)-3-[(3*S*,5*S*)-5-dimethylaminocarbonylpyrrolidin-3-ylthio]-6-[(*R*)-1-hydroxyethyl]-4-methyl-7-oxo-1-azabicyclo[3.2.0]hept-2-ene-2-carboxylic acid trihydrate, is confirmed. Meropenem crystallized as a zwitterion with the three molecules of water. The sum of the three bond angles about the N atom of the β -lactam ring is 329.1° and the deviation of the N atom from the plane defined by the three adjacent atoms is 0.457 \AA . Short intramolecular contacts are observed between the 1β -methyl group and the β -lactam ring.

Comment

Thienamycin and the related naturally occurring compounds are β -lactam antibiotics possessing potent antibacterial activity (Albers-Schönberg *et al.*, 1978). They are also chemically unstable and easily metabolized by renal dehydropeptidase-I (DHP-I). Meropenem (1) is a new 1β -methylcarbapenem antibiotic possessing excellent antibacterial activity and chemical stability with high stability to DHP-I (Sunagawa, Matsumura, Inoue, Fukasawa & Kato, 1990).



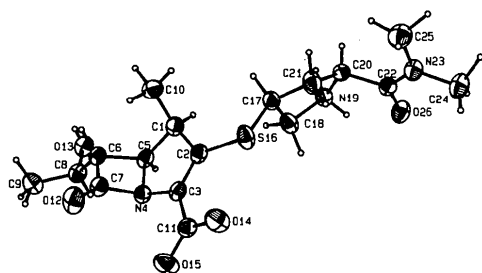


Fig. 1. View of meropenem showing the labelling of the non-H atoms. Thermal ellipsoids are shown at 50% probability levels. H atoms are drawn as small circles of arbitrary radius.

Meropenem crystallized as a zwitterion with a negative charge on the carboxyl group and a positive charge on the pyrrolidine ring N atom. The H atom H(N19A) of the pyrrolidine ring is separated by 2.00(9) Å from the O atom of the dimethylaminocarbonyl group, suggesting that there is an intramolecular hydrogen bond. The conformation of the dimethylaminocarbonyl group is fixed by the formation of an intramolecular hydrogen bond. Intermolecular hydrogen bonds are also found between the N atom of the pyrrolidine ring, the carboxyl group and the three water molecules.

The sum of the three bond angles around the β -lactam N atom (ΣN) is 329.1° and the distance of that N atom from the plane of its three attached C atoms (D) is 0.457 Å. These values are similar to those of the thienamycin derivative (Albers-Schönberg *et al.*, 1978) which has $\Sigma N = 325.9^\circ$ and $D = 0.49$ Å. Cohen (1983) evaluated the significance parameter for antibacterial activity by measuring the distance between the β -lactam O atom and the carboxylate C atom; distances between 3.0 and 3.9 Å were observed in active compounds. In the case of meropenem, the distance is 3.617(3) Å.

The 1 β -methyl group C(10) makes a short contact with the β -lactam ring. The steric effect of 1 β -methyl substitution may contribute to the enhanced chemical and metabolic stability of the meropenem.

Experimental

Crystal data

$C_{17}H_{25}N_3O_5 \cdot 3H_2O$

$M_r = 437.52$

Monoclinic

$P2_1$

$a = 9.279$ (1) Å

$b = 14.035$ (1) Å

$c = 9.123$ (1) Å

$\beta = 117.37$ (1)°

$V = 1055.1$ Å³

$Z = 2$

$D_x = 1.377$ Mg m⁻³

Data collection

Enraf-Nonius CAD-4 diffractometer

Cu $K\alpha$ radiation

$\lambda = 1.5418$ Å

Cell parameters from 25 reflections

$\theta = 21-29^\circ$

$\mu = 1.75$ mm⁻¹

$T = 295$ K

Prism

$0.45 \times 0.25 \times 0.05$ mm

Colorless

$\omega/2\theta$ scans

Absorption correction: none

2086 measured reflections

2086 independent reflections

1994 observed reflections

$[I > 3.0\sigma(I)]$

Refinement

Refinement on F

Final $R = 0.034$

$wR = 0.046$

$S = 1.820$

1994 reflections

386 parameters

All H-atom parameters refined

$w = 4/[\sigma^2(I) + 0.0025I^2]$

Data collection: Enraf-Nonius CAD-4 software. Cell refinement: Enraf-Nonius CAD-4 software. Data reduction: *BEGIN* (Enraf-Nonius, 1985). Program(s) used to solve structure: *MITHRIL* (Gilmore, 1984). Program(s) used to refine structure: *LSFM* (Enraf-Nonius, 1985). Molecular graphics: *ORTEP* (Johnson, 1976). Software used to prepare material for publication: *ESD*, *PTABLE* and *BTABLE* (Enraf-Nonius, 1985).

Table 1. Fractional atomic coordinates and equivalent isotropic thermal parameters (Å²)

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

	<i>x</i>	<i>y</i>	<i>z</i>	U_{eq}
S(16)	0.55458 (6)	0.5	0.74195 (6)	0.0399 (1)
O(12)	0.3509 (2)	0.7147 (2)	0.1656 (3)	0.0512 (6)
O(13)	0.9029 (2)	0.7615 (1)	0.3140 (2)	0.0388 (4)
O(14)	0.2714 (2)	0.4566 (2)	0.4370 (2)	0.0474 (5)
O(15)	0.2596 (2)	0.4789 (2)	0.1919 (2)	0.0527 (5)
O(26)	0.9236 (2)	0.2349 (1)	1.2017 (2)	0.0410 (4)
N(4)	0.5389 (2)	0.5967 (1)	0.3284 (2)	0.0293 (4)
N(19)	0.9733 (2)	0.3862 (2)	1.0718 (2)	0.0330 (5)
N(23)	0.7839 (2)	0.2972 (2)	1.3249 (2)	0.0348 (5)
C(1)	0.7161 (2)	0.6279 (2)	0.6129 (3)	0.0319 (5)
C(2)	0.5845 (2)	0.5536 (2)	0.5860 (3)	0.0299 (5)
C(3)	0.4799 (2)	0.5444 (2)	0.4245 (2)	0.0291 (5)
C(5)	0.7049 (2)	0.6340 (2)	0.4393 (3)	0.0294 (5)
C(6)	0.6578 (2)	0.7263 (2)	0.3333 (2)	0.0299 (5)
C(7)	0.4858 (2)	0.6852 (2)	0.2538 (3)	0.0324 (5)
C(8)	0.7379 (2)	0.7365 (2)	0.2200 (3)	0.0333 (5)
C(9)	0.6481 (3)	0.8061 (2)	0.0798 (3)	0.0450 (7)
C(10)	0.6765 (3)	0.7205 (2)	0.6755 (3)	0.0418 (6)
C(11)	0.3240 (2)	0.4899 (2)	0.3453 (3)	0.0320 (5)
C(17)	0.7558 (3)	0.4936 (2)	0.9184 (2)	0.0344 (5)
C(18)	0.8655 (3)	0.4192 (2)	0.8991 (3)	0.0385 (6)
C(20)	0.8868 (3)	0.4029 (2)	1.1749 (2)	0.0302 (5)
C(21)	0.7285 (3)	0.4534 (2)	1.0595 (3)	0.0401 (6)
C(22)	0.8653 (2)	0.3047 (2)	1.2365 (2)	0.0297 (5)
C(24)	0.7619 (3)	0.2022 (2)	1.3767 (3)	0.0448 (6)
C(25)	0.7156 (3)	0.3767 (2)	1.3742 (3)	0.0481 (7)
O(W1)	0.0866 (2)	0.4499 (2)	0.5999 (2)	0.0521 (6)
O(W2)	0.9964 (2)	0.1960 (2)	0.9142 (2)	0.0624 (7)
O(W3)	0.1787 (4)	0.5468 (2)	-0.1173 (3)	0.0848 (9)

Table 2. Geometric parameters (Å, °)

S(16)—C(2)	1.741 (3)	C(1)—C(2)	1.537 (3)
O(12)—C(7)	1.208 (3)	C(1)—C(5)	1.542 (4)
O(13)—C(8)	1.412 (2)	C(1)—C(10)	1.531 (4)
O(14)—C(11)	1.237 (3)	C(2)—C(3)	1.347 (3)
O(15)—C(11)	1.253 (3)	C(3)—C(11)	1.496 (3)
O(26)—C(22)	1.229 (3)	C(5)—C(6)	1.554 (3)

N(4)—C(3)	1.432 (3)	C(6)—C(7)	1.530 (3)
N(4)—C(5)	1.495 (2)	C(6)—C(8)	1.532 (4)
N(4)—C(7)	1.393 (3)	C(8)—C(9)	1.517 (3)
N(19)—C(18)	1.500 (3)	C(17)—C(18)	1.523 (4)
N(19)—C(20)	1.508 (4)	C(17)—C(21)	1.530 (4)
N(23)—C(22)	1.339 (4)	C(20)—C(21)	1.532 (3)
N(23)—C(24)	1.461 (4)	C(20)—C(22)	1.535 (3)
N(23)—C(25)	1.453 (4)	O(W1)—O(W3 ⁱⁱⁱ)	2.685 (3)
N(19)—O(26)	2.574 (3)	O(W2)—O(12 ^{iv})	2.971 (3)
N(19)—O(15 ⁱ)	2.697 (3)	O(W2)—O(13 ^v)	2.801 (3)
O(13)—O(W1 ⁱⁱ)	2.748 (3)	O(W3)—O(26 ⁱⁱ)	2.791 (3)
O(W1)—O(14)	2.739 (3)		
O(15)—O(W3)	2.735 (3)		
C(3)—N(4)—C(5)	109.3 (2)	C(5)—C(6)—C(8)	114.4 (2)
C(3)—N(4)—C(7)	127.4 (2)	C(7)—C(6)—C(8)	116.1 (2)
C(5)—N(4)—C(7)	92.4 (2)	O(12)—C(7)—N(4)	131.0 (2)
C(18)—N(19)—C(20)	108.5 (2)	O(12)—C(7)—C(6)	136.3 (2)
C(22)—N(23)—C(24)	117.9 (2)	N(4)—C(7)—C(6)	92.7 (2)
C(22)—N(23)—C(25)	125.0 (2)	O(13)—C(8)—C(6)	109.9 (2)
C(24)—N(23)—C(25)	117.1 (2)	O(13)—C(8)—C(9)	112.0 (2)
C(2)—C(1)—C(5)	101.6 (2)	C(6)—C(8)—C(9)	112.3 (2)
C(2)—C(1)—C(10)	109.1 (2)	O(14)—C(11)—O(15)	125.1 (2)
C(5)—C(1)—C(10)	114.9 (2)	O(14)—C(11)—C(3)	117.4 (2)
S(16)—C(2)—C(1)	125.1 (1)	O(15)—C(11)—C(3)	117.4 (2)
S(16)—C(2)—C(3)	123.2 (2)	C(18)—C(17)—C(21)	102.8 (2)
C(1)—C(2)—C(3)	111.1 (2)	N(19)—C(18)—C(17)	104.4 (2)
N(4)—C(3)—C(2)	109.5 (2)	N(19)—C(20)—C(21)	105.1 (2)
N(4)—C(3)—C(11)	121.6 (2)	N(19)—C(20)—C(22)	106.5 (2)
C(2)—C(3)—C(11)	128.9 (2)	C(21)—C(20)—C(22)	114.9 (2)
N(4)—C(5)—C(1)	103.8 (2)	C(17)—C(21)—C(20)	105.7 (2)
N(4)—C(5)—C(6)	87.9 (1)	O(26)—C(22)—N(23)	122.0 (2)
C(1)—C(5)—C(6)	123.6 (2)	O(26)—C(22)—C(20)	118.3 (2)
C(5)—C(6)—C(7)	85.1 (2)	N(23)—C(22)—C(20)	119.7 (2)

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

The ω -scan width was $(1.0 + 0.15 \tan \theta)^\circ$. Refinement was by full-matrix least-squares methods. Refinement of the other enantiomer under identical conditions yielded $R = 0.038$, $wR = 0.053$ and $S = 2.077$. The better fit for the atomic coordinates presented in Table 1 is in agreement with the known absolute configuration. Calculations were performed on a MicroVAX II computer.

Lists of structure factors, anisotropic thermal parameters, H-atom coordinates and complete geometry have been deposited with the British Library Document Supply Centre as Supplementary Publication No. SUP 55253 (23 pp.). Copies may be obtained through The Technical Editor, International Union of Crystallography, 5 Abbey Square, Chester CH1 2HU, England. [CIF reference: AS1005]

References

- Albers-Schönberg, G., Arison, B. H., Hensens, O. D., Hirshfield, J., Hoogsteen, K., Kaczka, E. A., Rhodes, R. E., Kahan, J. S., Kahan, F. M., Ratcliffe, R. W., Walton, E., Ruswinkle, L. J., Morin, R. B. & Christensen, B. G. (1978). *J. Am. Chem. Soc.* **100**, 6491–6499.
- Cohen, N. C. (1983). *J. Med. Chem.* **26**, 259–264.
- Enraf-Nonius (1985). *Structure Determination Package*. Enraf-Nonius, Delft, The Netherlands.
- Gilmore, C. J. (1984). *J. Appl. Cryst.* **17**, 42–46.
- Johnson, C. K. (1976). *ORTEPII*. Report ORNL-5138. Oak Ridge National Laboratory, Tennessee, USA.
- Sunagawa, M., Matsumura, H., Inoue, T., Fukasawa, M. & Kato, M. (1990). *J. Antibiot.* **43**, 519–532.