

Table I. Crystal Parameters for Antibiotic A23187

<i>a</i>	15.759 (4) Å
<i>b</i>	10.377 (4) Å
<i>c</i>	8.592 (3) Å
β	95.97 (2)°
Space group	$P2_1$
Molecules/cell	2
Obsd density	1.264 g cm ⁻³
Calcd density	1.244 g cm ⁻³

The structure was solved by direct methods using the program MULTAN³ and was refined by the least-squares method. The final R factor, using anisotropic temperature factors for all heavy atoms and isotropic temperature factors for all hydrogen atoms (at assumed locations), was 0.063. The refined atomic coordinates for the heavy atoms are given in Table II.⁴ The conformation of the molecule in the crystal is shown in Figure 1.

The structure consists of three basic units, α -keto-pyrrole, a substituted benzoxazole, and a spiro ring system similar to those found in the polyether antibiotics—monensin,⁵ nigericin,⁶ grisorixin,⁷ dianemycin,⁸ X-206,⁹ and A204A.¹⁰ In the polyether antibiotics, the spiro ring systems consist of a five- and a six-membered ring, whereas A23187 contains two six-membered rings. Comparison of the spiro moieties in the polyethers and in A23187 shows that conformationally, they are very similar; in each case, the ring ether oxygen atoms are in axial or pseudoaxial conformations, and the points of attachment of the rest of the molecular chain are equatorial or pseudoequatorial.

Because the molecule contains no atom with strong anomalous X-ray scattering, we have been unable to determine experimentally the absolute configuration. However, in Figure 1, the chiralities of the asymmetric centers in each of the two six-membered rings of the spiro group are the same as those found in all the polyethers which contain a spiro six-membered ring.¹¹ It seems probable, therefore, that the absolute configuration shown is correct.

In the crystalline, free acid form of A23187, there are three internal hydrogen bonds, as shown by the dotted lines in Figure 1. The hydrogen bond between the pyrrole nitrogen atom and one of the carboxyl oxygen atoms holds the ends of the molecule in close proximity,

(3) P. Main, M. M. Woofson, and G. Germain, "MULTAN, a Computer Programme for the Automatic Solution of Crystal Structures," University of York Printing Unit, York, England, 1971.

(4) See paragraph at end of paper regarding supplementary material.

(5) A. Agtarap, J. W. Chamberlin, M. Pinkerton, and L. Steinrauf, *J. Amer. Chem. Soc.*, **89**, 5737 (1967); M. Pinkerton and L. K. Steinrauf, *J. Mol. Biol.*, **49**, 533 (1970); W. K. Lutz, F. K. Winkler, and J. D. Dunitz, *Helv. Chim. Acta*, **54**, 1103 (1971).

(6) L. K. Steinrauf, M. Pinkerton, and J. W. Chamberlin, *Biochem. Res. Commun.*, **33**, 29 (1968); T. Kubota, S. Matsutani, M. Shiro, and H. Koyama, *Chem. Commun.*, 1541 (1968); T. Kubota and S. Matsutani *J. Chem. Soc. C*, 695 (1970).

(7) P. Gachon, A. Kergomard, H. Veschambre, C. Esteve, and T. Staron, *Chem. Commun.*, 1421 (1970); M. Alleaume and D. Hickel, *ibid.*, 1422 (1970); M. Alleaume and D. Hickel, *J. Chem. Soc., Chem. Commun.*, 175 (1972).

(8) R. L. Hamill, M. M. Hoehn, G. E. Pittenger, J. Chamberlin, and M. Gorman, *J. Antibiot.*, **22**, 161 (1969); E. W. Czerwinski and L. K. Steinrauf, *Biochem. Biophys. Res. Commun.*, **45**, 1284 (1971).

(9) J. F. Blount and J. W. Westley, *Chem. Commun.*, 927 (1971).

(10) N. D. Jones, M. O. Chaney, J. W. Chamberlin, R. L. Hamill, and S. Chen, *J. Amer. Chem. Soc.*, **95**, 3399 (1973).

(11) Dianemycin (ref 8) contains two spiro ring systems. The one closest to the carboxyl end of the molecule has the same chiralities found in the polyether antibiotics mentioned above, but the other spiro moiety has several asymmetric centers with reversed chirality.

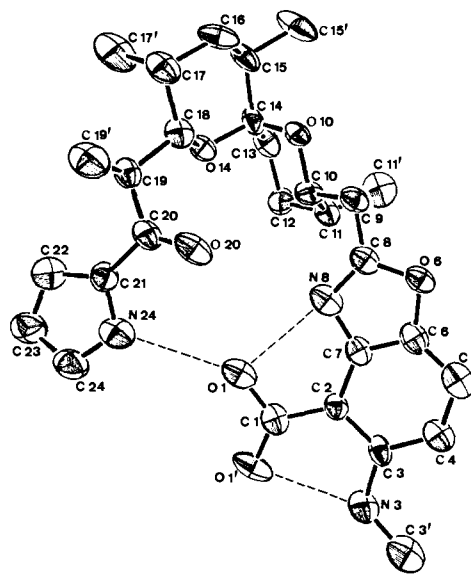


Figure 1. Conformation and proposed absolute configuration of A23187 in the crystal. The thermal ellipsoids are drawn at the 50% probability level.

a feature usually seen in the crystal structures of polyether antibiotics. Although the structure of the 2:1 antibiotic-divalent cation complex is not yet known, one can speculate that the principal ligands to the metal ion are the oxygen atoms of the carbonyl and carboxyl group, as well as one of the ether oxygen atoms from the spiro ring system.

Acknowledgment. We wish to thank Dr. R. L. Hamill and Mr. E. A. Presti for supplying the antibiotic used in this investigation and Dr. D. E. Dorman for his help in the preparation of this manuscript. Appreciation is also due Mr. D. W. Smith for computer assistance and Mr. J. W. Paschal and Mr. J. P. Hettler for their help in obtaining nmr and mass spectra.

Supplementary Material Available. A listing of refined atomic coordinates will appear following these pages in the microfilm edition of this volume of the journal. Photocopies of the supplementary material from this paper only or microfiche (105 × 148 mm, 24× reduction, negatives) containing all of the supplementary material for the papers in this issue may be obtained from the Journals Department, American Chemical Society, 1155 16th St., N.W., Washington, D. C. 20036. Remit check or money order for \$3.00 for photocopy or \$2.00 for microfiche, referring to code number JACS-74-1932.

Michael O. Chaney,* Paul V. Demarco
Noel D. Jones, John L. Occolowitz

The Lilly Research Laboratories, Eli Lilly and Company
Indianapolis, Indiana 46206

Received November 9, 1973

Rapid Relaxation of Spin Equilibrium in Ferric Myoglobin Hydroxide

Sir:

The hypothesis¹ that changes in the spin states of the cytochromes are intimately related to the mechanism of oxidative phosphorylation has attractive features. Electron transfer and phosphorylation could be coupled through the conformational changes induced in the

(1) D. F. Wilson, P. L. Dutton, M. Erecinska, J. G. Lindsay, and N. Sato, *Accounts Chem. Res.*, **5**, 234 (1972), and references therein.