

parallel circular layers subject to rotational and displacement disordering.

1. Using the Gaussian approximation for the interference functions obtained (equation (10)) the 002 reflections for microcrystalline carbon were investigated.

- (a) In the absence of displacement disorder, use of the tangent plane approximation for interpreting the 002 diffraction maximum results in a small systematic error in evaluating the interplanar spacing, a_3 . For a layer radius in excess of 5 Å, the value of a_3 using the *TPA* will be too small by less than 2%.
- (b) While rotational disorder plays a negligible rôle in the 002 diffraction, displacement disorder can seriously affect the peak profile. As the displacement disorder increases, the 002 intensity diminishes toward zero. The effect of one type of displacement disorder is evaluated. It is found that experimentally one would find a negligible difference between a given number of crystals showing no displacement disorder and a larger number of crystals showing appreciable displacement disorder. Thus experimental estimates of the fraction of layers stacked in parallel clusters will be a sensitive function of the displacement disorder present in those clusters.
- (c) Expressions for small angle scattering (000 Bragg maximum) can be deduced directly from ex-

pressions for the 002 profile because the *TPA* was not used. The low angle equations give useful approximations to results obtained by numerical integration of the actual interference functions.

2. Results of a similar analysis on the $hk0$ maxima showed that random rotational disorder by itself effectively suppresses interlayer interference effects in carbon for $R > 5$ Å.

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Short Communications

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On the crystal structure of aureomycin hydrochloride, By S. HIROKAWA, Y. OKAYA, F. M. LOVELL and R. PEPINSKY, *X-ray and Crystal Structure Laboratory, Department of Physics, The Pennsylvania State University, University Park, Pa., U.S.A.*

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A structure analysis of aureomycin hydrochloride, $C_{22}H_{23}N_2O_8Cl \cdot HCl$, was undertaken in order to clarify certain stereochemical features of this member of the tetracyclin family. The salt crystallizes in the orthorhombic system with four chemical units in the cell, the dimensions of which are

$$a = 11.20, \quad b = 12.89, \quad c = 15.47 \text{ \AA},$$

with space group $P2_12_12_1$ (Dunitz & Leonard, 1950; Pepinsky & Watanabé, 1952). Crystalline terramycin hydrochloride is strikingly isomorphous (Pepinsky & Watanabé, 1952).

X-ray data were collected as multi-film Weissenberg exposures, using $Cu K\alpha$ radiation, with intensities visually estimated. The structure analysis was initiated by establishment of the chlorine positions through a sys-

tematic study of a three-dimensional sharpened Patterson function. Starting from these positions, numerous iterated structure-factor and electron-density calculations on the IBM 704 and X-RAC were made, with contributions of lighter atoms included as they became discernible in the electron-density maps. Interpretations of these were aided by bounded projections along the a and b axes. All the atomic coordinates thus obtained were subjected first to least-squares and then differential-synthesis refinement, using the automatic refinement program of Vand & Pepinsky (1958) on the IBM 704. A three-dimensional error index of 0.154 was obtained for the final set of coordinates.

The aureomycin ion is found to possess the structure as proposed on chemical grounds (Hochstein *et al.*, 1953), with the exception of one significant stereochemical

feature and various details of bond character and probable hydrogen locations. The dimethylamino group on C(4) takes the polar (ϵ) configuration with respect to ring I to which it is attached. This is the converse of the configuration assigned chemically (Woodward, 1959), and the X-ray structure thus corresponds to the conformation assigned on chemical grounds to *epi*-aureomycin. The terminology for these epimers has been re-examined through the cooperation of Dr J. H. Boothe of the Lederle Laboratories, who submitted fresh samples of the hydrochlorides of aureomycin and *epi*-aureomycin. Powder and single-crystal examination of these confirmed the fact that the structure we report here is that of aureomycin, HCl.

Some features of the stereochemistry of the aureomycin ion are given in Fig. 1(a), and bond distances are also given in this figure. [Standard deviations of coordinates have been assessed by the method of Cruickshank (1949), and are in the neighborhood of 0.02 Å.] The various atoms of the ion are shown in Fig. 1(b), as first numbered by Hochstein *et al.* (1953). Dashed lines indicate bonds below the plane of Fig. 1(a). Several distances indicating intramolecular hydrogen bonds are found. These are indicated by dotted lines. It is interesting that O(11) appears to be involved in hydrogen bonding to both O(10) and O(12); and both C(10)–O(10) and C(12)–O(12) appear to be of double-bond (ketonic) character, while C(11)–O(11) appears to be a single bond (carbon to hydroxyl). This is in disagreement with earlier chemically-derived conclusions (Hochstein *et al.*, 1953). Aromatic ring IV has a partially quinone character. The carbon sequences C(10a)–C(11)–C(11a)–C(12) and C(1)–C(2)–C(3) are conjugated. Distances from C_{amide}(2) to the oxygen and nitrogen of the amide group are anomalous, in that the C–N and C–O separations are opposite to those found

in formamide (Ladell & Post, 1954), oxamide (Ayerst & Duke, 1954) and nicotinamide (Wright & King, 1954); but the identity of N and O here is substantiated both by the electron-density maps and (very convincingly) by the distances N_{amide}(2)–O(1) [= 2.80 Å] and O_{amide}(2)–O(3) [= 2.51 Å]. The character of these bonds can be explained by an examination of the resonance state of the molecule, the most important effect on which comes from the influence of the dimethylamino group on the basicity of the amide group. This problem is considered elsewhere, in a complete report of the analysis (Hirokawa *et al.*, 1959).

The chloride ion is surrounded by six atoms in a distorted octahedron, contributed from three symmetry-related aureomycin ions. These atoms are N(4), O(6) and CH₃(6) of one ion, at distances 3.10, 3.22 and 3.72 Å, respectively; O(12) and O(12a) of another ion, at distances 3.31 and 3.22 Å, respectively; and N_{amide}(2) of a third ion, at 3.12 Å. Aside from hydrogen bonding to this chloride ion, all other intermolecular contacts are of van der Waals type.

Full details of this analysis, and discussions of various chemical features of the structure, are in press in the report referred to above (Hirokawa *et al.*, 1959).

Since terramycin hydrochloride is strikingly isomorphous with aureomycin hydrochloride, a refinement of the former structure is now in progress, utilizing atomic positions as found in aureomycin but of course omitting Cl(7) and adding a hydroxyl at C(5).

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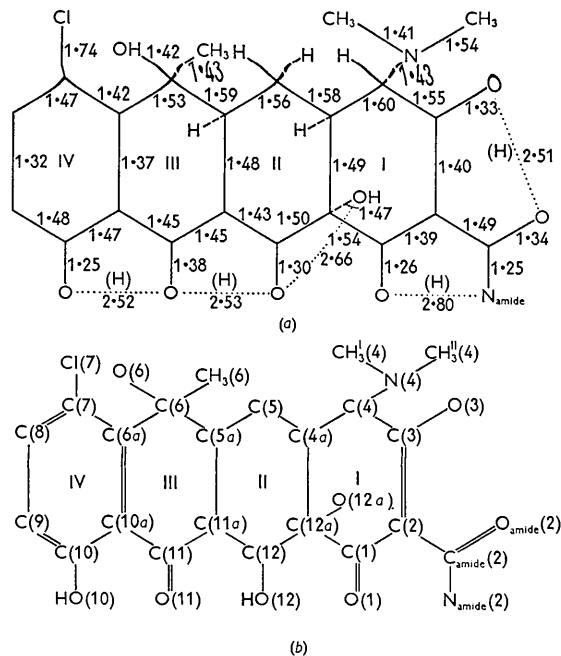


Fig. 1. (a) Aureomycin, HCl: Some intra-molecular bond distances as found from X-ray analysis. Dashed lines indicate bonds below plane of figure; dotted lines indicate hydrogen bonds. (b) Aureomycin: Molecular formula and numbering as given by Hochstein *et al.* (1953).