

Crystal Structure of Methyl α -D-Mannopyranoside

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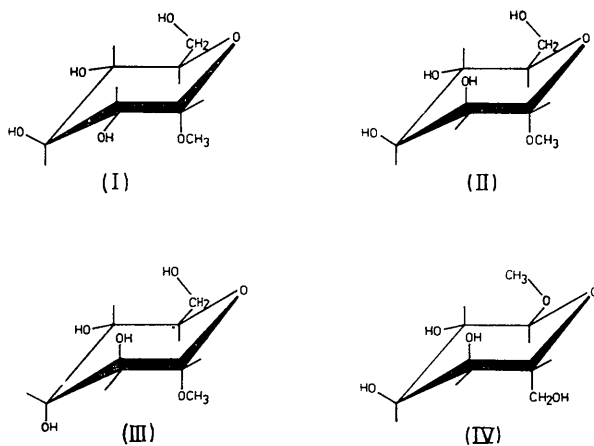
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Methyl α -D-mannopyranoside crystallizes in the orthorhombic system: space group $P2_12_12_1$. Cell dimensions at room temperature are $a=9.42_3$, $b=9.30_8$ and $c=10.04_5$ Å. The structure was solved by direct phasing methods with automatic diffractometer data and refined to a final reliability index of 0.044. Bond lengths and angles are similar to those encountered in other pyranose sugars and all oxygen atoms are involved in the hydrogen bonding network which comprises two terminating chains.

Introduction

The methyl α -glycosides of glucose, mannose and altrose constitute a valuable series of compounds in the study of hydrogen bonding patterns and conformational stability in pyranose sugars and their derivatives.

In changing from methyl α -glucoside (I) through methyl α -mannoside (II) to methyl α -altroside (III) the number of axial substituents in the $C1$ conformation is increased from one to three. At the same time the energy difference of the two alternative chair conformations for each compound decreases as steric interaction between the more bulky hydroxyl axial groups increases. With methyl α -altroside this results in the energies of the two conformations (III, IV) being almost identical (Eliel, Allinger, Angyal & Morrison, 1965) so that the conformation of the molecule in the crystal will be controlled by hydrogen bonding and packing requirements. Interest in this series of compounds is heightened further by the similar configuration of $C1$ methyl α -mannoside (II) and $1C$ methyl α -altroside (IV). All three materials crystallize in the orthorhombic space group $P2_12_12_1$ with a single molecule per asymmetric unit.



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The structure of methyl α -glucoside has been determined (Berman & Kim, 1968): this paper reports that of methyl α -mannoside.

Experimental

Methyl α -D-mannopyranoside, $C_7H_{14}O_6$, F.W. 194.19
Orthorhombic:

$$\begin{aligned} a &= 9.42_3 \text{ \AA} \\ b &= 9.30_8 \\ c &= 10.04_5 \\ D_m &= 1.459 \text{ g.cm}^{-3} \quad D_x = 1.463 \text{ g.cm}^{-3} \end{aligned}$$

Systematic absences: $h00$ for h odd; $0k0$ for k odd; $00l$ for l odd

Space group: $P2_12_12_1$

$$\begin{aligned} \mu &11.29 \text{ cm}^{-1} \\ \text{Cu } K\alpha_1 &1.5405 \text{ \AA} \end{aligned}$$

These cell dimensions were measured with a Picker automatic four-circle diffractometer using high-order axial reflexions for which the $\text{Cu } K\alpha_1$ and α_2 peaks were resolved. A set was also obtained from film data with Si ($a=5.4305$ Å) as internal standard. The agreement between the two sets was better than 0.25% but both were approximately 1% greater than those recorded by Cox & Goodwin (1932).

Three sets of intensity data were collected. Two were from different octants of the same crystal using the diffractometer operating in the $\theta-2\theta$ scanning mode with a scan rate of $0.5^\circ \text{ min}^{-1}$ and stationary-crystal stationary-counter background counts of 40 sec at each end of the scan. The third set comprised film data collected at room temperature about two axes and cross-correlated *via* common reflexions. Intensities were estimated visually on a logarithmic scale against an appropriate standard series.

The common observed structure factors of the two diffractometer data sets had an agreement index, $R_f = 2\sum||F_1| - |F_2|| / \sum(|F_1| + |F_2|)$, of 0.015; the corresponding R_f between diffractometer and film data was 0.065. The diffractometer datum set used in the refinement was chosen arbitrarily.

All crystals were approximate cubes with edges 0.15–0.20 mm, obtained by recrystallization from ethanol of material supplied by Nutritional Biochemical Corporation. Lorentz-polarization corrections were applied to all intensities but no allowance was made for absorption. The structure factors were placed on an approximate absolute scale using a Wilson plot and final scaling was achieved by equating the calculated to the observed values of the structure factors during the least-squares refinement.

The atomic scattering factors for carbon, oxygen, and hydrogen were those of Freeman (1959), Hoerni & Ibers (1954) and Stewart, Davidson & Simpson (1965) respectively.

Structure determination and refinement

The initial attempt to solve the structure was made using a vector map synthesized with film data. Several plausible orientations of the molecule for both chair conformations (*C*1 and *1C*), were deduced from the 2.4 Å vector distribution about the origin but numerous attempts to locate the molecule within the cell using Harker sections were unsuccessful.

The positioning of the molecule was next approached by generating an 'R' map for a particular molecular orientation and conformation. This *R* map was produced by calculating the conventional reliability index, *R*, for seventeen low-angle reflexions as the molecular model was stepped systematically through the asymme-

tric unit. A number of minima were found but comparison of the observed and calculated structure factors at each position showed poor agreement which could not be improved satisfactorily using Fourier and difference syntheses. The probable reasons for failure were: firstly, most of the seventeen inner reflexions suffered significantly from extinction and/or absorption; secondly, only part of the molecule could be specified, *i.e.* the labile primary hydroxyl group and the methyl group were omitted from the model. Of the two factors the second was undoubtedly the more important.

The structure was finally solved by direct phasing methods using programs* incorporating the tangent formula of Karle & Hauptman (1956). Initial phases were assigned to three origin-defining reflexions of high *E* value and three structure invariant phases, the latter using the *E*₁ formula. The repeated application of the tangent formula with decreasing *E* values yielded 123 phased reflexions with *E* ≥ 1.53. From the resultant *E* map eleven atoms of the molecule were clearly revealed with some indication of the remaining two. An electron density map phased on these eleven confirmed the location of all thirteen heavy atoms; a knowledge of the expected structure of methyl mannose allowed the carbon and oxygen atoms to be differentiated from the beginning.

* A modified set of the D.P. suite of programs written by Dr S. R. Hall.

Table 1. Fractional atomic coordinates and anisotropic thermal parameters for methyl α -D-mannopyranoside

The temperature factor expression is

$$\exp [-(h^2\beta_{11} + k^2\beta_{22} + l^2\beta_{33} + 2hk\beta_{12} + 2hl\beta_{13} + 2kl\beta_{23})].$$

Estimated standard deviations are given in parentheses and refer to the last significant figure. $\beta_{ij} \times 10^4$.

| | <i>x/a</i> | <i>y/b</i> | <i>z/c</i> | β_{11} | β_{22} | β_{33} | β_{12} | β_{13} | β_{23} |
|----------|------------|-------------|------------|--------------|--------------|--------------|--------------|--------------|--------------|
| C(1) | 0.3591 (5) | 0.0419 (5) | 0.3896 (5) | 51 (5) | 49 (5) | 47 (5) | -2 (5) | -2 (5) | -11 (5) |
| C(2) | 0.4749 (5) | 0.0738 (5) | 0.2887 (5) | 56 (6) | 41 (5) | 43 (5) | 1 (5) | -5 (5) | 3 (5) |
| C(3) | 0.6205 (5) | 0.0490 (5) | 0.3489 (5) | 49 (6) | 39 (5) | 37 (5) | 2 (5) | 1 (5) | -2 (5) |
| C(4) | 0.6369 (5) | 0.1324 (5) | 0.4784 (5) | 49 (6) | 48 (5) | 38 (5) | 2 (5) | -4 (5) | 9 (5) |
| C(5) | 0.5158 (5) | 0.0960 (5) | 0.5726 (5) | 50 (6) | 60 (6) | 36 (5) | 4 (5) | 3 (5) | 7 (5) |
| C(6) | 0.5153 (7) | 0.1866 (6) | 0.6983 (5) | 93 (8) | 95 (7) | 41 (5) | 7 (7) | -1 (6) | -27 (6) |
| C(7) | 0.2571 (7) | -0.1561 (7) | 0.5053 (6) | 108 (8) | 113 (9) | 74 (7) | -40 (7) | 36 (7) | 2 (7) |
| O(1) | 0.3644 (4) | -0.1069 (3) | 0.4135 (3) | 84 (5) | 46 (4) | 56 (4) | -17 (4) | 25 (4) | -1 (4) |
| O(2) | 0.4546 (4) | 0.2217 (4) | 0.2528 (4) | 60 (5) | 51 (4) | 75 (5) | 4 (4) | -3 (4) | 28 (4) |
| O(3) | 0.7284 (4) | 0.0974 (4) | 0.2585 (3) | 55 (4) | 72 (5) | 42 (4) | -10 (4) | 12 (4) | -8 (4) |
| O(4) | 0.7672 (3) | 0.0891 (3) | 0.5429 (3) | 44 (4) | 66 (5) | 50 (4) | -9 (4) | -1 (3) | 7 (3) |
| O(5) | 0.3807 (3) | 0.1223 (3) | 0.5081 (3) | 49 (4) | 60 (4) | 46 (3) | 5 (4) | -1 (3) | -16 (3) |
| O(6) | 0.5133 (4) | 0.3360 (4) | 0.6715 (4) | 58 (4) | 87 (5) | 104 (4) | 3 (4) | -2 (4) | -44 (4) |
| HC(1) | 0.258 (5) | 0.068 (5) | 0.357 (4) | | | | | | |
| HC(2) | 0.462 (5) | 0.003 (5) | 0.194 (4) | | | | | | |
| HC(3) | 0.633 (5) | -0.048 (5) | 0.368 (4) | | | | | | |
| HC(4) | 0.633 (5) | 0.248 (5) | 0.456 (4) | | | | | | |
| HC(5) | 0.528 (4) | -0.019 (5) | 0.589 (4) | | | | | | |
| HC(6') | 0.425 (5) | 0.156 (5) | 0.763 (4) | | | | | | |
| HC(6'') | 0.622 (5) | 0.167 (5) | 0.748 (4) | | | | | | |
| HC(7') | 0.177 (5) | -0.115 (5) | 0.481 (5) | | | | | | |
| HC(7'') | 0.248 (5) | -0.261 (5) | 0.489 (5) | | | | | | |
| HC(7''') | 0.294 (5) | -0.131 (5) | 0.600 (5) | | | | | | |
| HO(2) | 0.507 (6) | 0.249 (6) | 0.215 (5) | | | | | | |
| HO(3) | 0.774 (6) | -0.048 (5) | 0.694 (5) | | | | | | |
| HO(4) | 0.833 (6) | 0.167 (7) | 0.538 (5) | | | | | | |
| HO(6) | 0.428 (6) | 0.365 (6) | 0.006 (5) | | | | | | |

Isotropic temperature factor = 2.2 Å

The structure was refined by full-matrix least-squares (Busing, Martin & Levy, 1962) using $\sum w(F_o - F_c)^2$. The film data were refined with both unit and individual weights. The latter were set proportional to $1/F^2$ which was appropriate to the method used in estimating intensity. Both schemes yielded a final overall R of 0.095. With unit weights a satisfactory agreement, R , with $\sin\theta/\lambda$ was obtained in that each zone was equally reliable; with individual weights there was significantly poorer agreement at higher angles. The atomic positional parameters were altered by up to 10σ in changing the weighting scheme but the unit weight refined parameters were within 3σ of the final diffractometer refined values. A difference electron density synthesis with $\sin\theta/\lambda \leq 0.45$ ($R=0.095$,

unit weights) showed peaks at acceptable hydrogen atom locations. However numerous other peaks of equal height were also present which vitiated the confident assignment of any of the peaks to hydrogen.

Refinement was continued using diffractometer data with individual weights based on counting statistics. A difference electron synthesis ($R=0.074$; $\sin\theta/\lambda \leq 0.45$) revealed most of the hydrogen atoms. Those attached to carbon atoms, with the exception of the methyl group, were well defined and with continued refinement all hydrogen atoms were eventually located unambiguously. The final difference Fourier synthesis showed electron density fluctuations not exceeding $0.4 \text{ e.}\text{\AA}^{-3}$ ($R=0.044$). Several low-angle reflexions showed evidence of absorption or extinction but in this refinement no reflexions were rejected.

Final values of the atomic coordinates and thermal parameters, together with their estimated standard deviations are given in Table 1; Table 2 contains the list of observed and calculated structure factors.

Description of the structure

Methyl α -D-mannopyranoside in the crystal has the C1 chair conformation for which the ring substituents are arranged in the sequence $1a2a3e4e5e$. The molecule and the numbering of the atoms is shown in Fig. 1.

Bond lengths and angles derived from the coordinates of Table 1 are listed in Table 3. The carbon-carbon bonds are between 1.517 and 1.524 Å with a mean of 1.520 Å which is in good agreement with values reported for other carbohydrates. The carbon-oxygen bond distances, however, show a greater variation and range between 1.407 and 1.450 Å with a mean value of 1.432 Å; this mean is not significantly greater

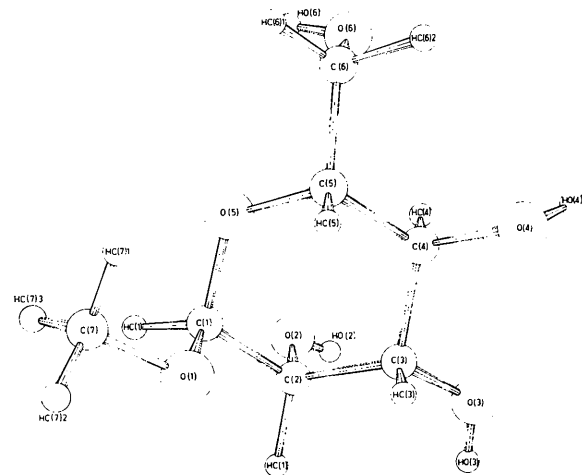


Fig. 1. Conformation of α -D-mannopyranoside and atom numbering.

Table 2. Calculated and observed structure factors

| h | k | l | F _o | F _c | ΔF | σ(F _o) | σ(F _c) |
|---|---|-----|----------------|----------------|----|--------------------|--------------------|
| 0 | 0 | 0 | 100 | 100 | 0 | 0 | 0 |
| 0 | 0 | 1 | 100 | 100 | 0 | 0 | 0 |
| 0 | 0 | 2 | 100 | 100 | 0 | 0 | 0 |
| 0 | 0 | 3 | 100 | 100 | 0 | 0 | 0 |
| 0 | 0 | 4 | 100 | 100 | 0 | 0 | 0 |
| 0 | 0 | 5 | 100 | 100 | 0 | 0 | 0 |
| 0 | 0 | 6 | 100 | 100 | 0 | 0 | 0 |
| 0 | 0 | 7 | 100 | 100 | 0 | 0 | 0 |
| 0 | 0 | 8 | 100 | 100 | 0 | 0 | 0 |
| 0 | 0 | 9 | 100 | 100 | 0 | 0 | 0 |
| 0 | 0 | 10 | 100 | 100 | 0 | 0 | 0 |
| 0 | 0 | 11 | 100 | 100 | 0 | 0 | 0 |
| 0 | 0 | 12 | 100 | 100 | 0 | 0 | 0 |
| 0 | 0 | 13 | 100 | 100 | 0 | 0 | 0 |
| 0 | 0 | 14 | 100 | 100 | 0 | 0 | 0 |
| 0 | 0 | 15 | 100 | 100 | 0 | 0 | 0 |
| 0 | 0 | 16 | 100 | 100 | 0 | 0 | 0 |
| 0 | 0 | 17 | 100 | 100 | 0 | 0 | 0 |
| 0 | 0 | 18 | 100 | 100 | 0 | 0 | 0 |
| 0 | 0 | 19 | 100 | 100 | 0 | 0 | 0 |
| 0 | 0 | 20 | 100 | 100 | 0 | 0 | 0 |
| 0 | 0 | 21 | 100 | 100 | 0 | 0 | 0 |
| 0 | 0 | 22 | 100 | 100 | 0 | 0 | 0 |
| 0 | 0 | 23 | 100 | 100 | 0 | 0 | 0 |
| 0 | 0 | 24 | 100 | 100 | 0 | 0 | 0 |
| 0 | 0 | 25 | 100 | 100 | 0 | 0 | 0 |
| 0 | 0 | 26 | 100 | 100 | 0 | 0 | 0 |
| 0 | 0 | 27 | 100 | 100 | 0 | 0 | 0 |
| 0 | 0 | 28 | 100 | 100 | 0 | 0 | 0 |
| 0 | 0 | 29 | 100 | 100 | 0 | 0 | 0 |
| 0 | 0 | 30 | 100 | 100 | 0 | 0 | 0 |
| 0 | 0 | 31 | 100 | 100 | 0 | 0 | 0 |
| 0 | 0 | 32 | 100 | 100 | 0 | 0 | 0 |
| 0 | 0 | 33 | 100 | 100 | 0 | 0 | 0 |
| 0 | 0 | 34 | 100 | 100 | 0 | 0 | 0 |
| 0 | 0 | 35 | 100 | 100 | 0 | 0 | 0 |
| 0 | 0 | 36 | 100 | 100 | 0 | 0 | 0 |
| 0 | 0 | 37 | 100 | 100 | 0 | 0 | 0 |
| 0 | 0 | 38 | 100 | 100 | 0 | 0 | 0 |
| 0 | 0 | 39 | 100 | 100 | 0 | 0 | 0 |
| 0 | 0 | 40 | 100 | 100 | 0 | 0 | 0 |
| 0 | 0 | 41 | 100 | 100 | 0 | 0 | 0 |
| 0 | 0 | 42 | 100 | 100 | 0 | 0 | 0 |
| 0 | 0 | 43 | 100 | 100 | 0 | 0 | 0 |
| 0 | 0 | 44 | 100 | 100 | 0 | 0 | 0 |
| 0 | 0 | 45 | 100 | 100 | 0 | 0 | 0 |
| 0 | 0 | 46 | 100 | 100 | 0 | 0 | 0 |
| 0 | 0 | 47 | 100 | 100 | 0 | 0 | 0 |
| 0 | 0 | 48 | 100 | 100 | 0 | 0 | 0 |
| 0 | 0 | 49 | 100 | 100 | 0 | 0 | 0 |
| 0 | 0 | 50 | 100 | 100 | 0 | 0 | 0 |
| 0 | 0 | 51 | 100 | 100 | 0 | 0 | 0 |
| 0 | 0 | 52 | 100 | 100 | 0 | 0 | 0 |
| 0 | 0 | 53 | 100 | 100 | 0 | 0 | 0 |
| 0 | 0 | 54 | 100 | 100 | 0 | 0 | 0 |
| 0 | 0 | 55 | 100 | 100 | 0 | 0 | 0 |
| 0 | 0 | 56 | 100 | 100 | 0 | 0 | 0 |
| 0 | 0 | 57 | 100 | 100 | 0 | 0 | 0 |
| 0 | 0 | 58 | 100 | 100 | 0 | 0 | 0 |
| 0 | 0 | 59 | 100 | 100 | 0 | 0 | 0 |
| 0 | 0 | 60 | 100 | 100 | 0 | 0 | 0 |
| 0 | 0 | 61 | 100 | 100 | 0 | 0 | 0 |
| 0 | 0 | 62 | 100 | 100 | 0 | 0 | 0 |
| 0 | 0 | 63 | 100 | 100 | 0 | 0 | 0 |
| 0 | 0 | 64 | 100 | 100 | 0 | 0 | 0 |
| 0 | 0 | 65 | 100 | 100 | 0 | 0 | 0 |
| 0 | 0 | 66 | 100 | 100 | 0 | 0 | 0 |
| 0 | 0 | 67 | 100 | 100 | 0 | 0 | 0 |
| 0 | 0 | 68 | 100 | 100 | 0 | 0 | 0 |
| 0 | 0 | 69 | 100 | 100 | 0 | 0 | 0 |
| 0 | 0 | 70 | 100 | 100 | 0 | 0 | 0 |
| 0 | 0 | 71 | 100 | 100 | 0 | 0 | 0 |
| 0 | 0 | 72 | 100 | 100 | 0 | 0 | 0 |
| 0 | 0 | 73 | 100 | 100 | 0 | 0 | 0 |
| 0 | 0 | 74 | 100 | 100 | 0 | 0 | 0 |
| 0 | 0 | 75 | 100 | 100 | 0 | 0 | 0 |
| 0 | 0 | 76 | 100 | 100 | 0 | 0 | 0 |
| 0 | 0 | 77 | 100 | 100 | 0 | 0 | 0 |
| 0 | 0 | 78 | 100 | 100 | 0 | 0 | 0 |
| 0 | 0 | 79 | 100 | 100 | 0 | 0 | 0 |
| 0 | 0 | 80 | 100 | 100 | 0 | 0 | 0 |
| 0 | 0 | 81 | 100 | 100 | 0 | 0 | 0 |
| 0 | 0 | 82 | 100 | 100 | 0 | 0 | 0 |
| 0 | 0 | 83 | 100 | 100 | 0 | 0 | 0 |
| 0 | 0 | 84 | 100 | 100 | 0 | 0 | 0 |
| 0 | 0 | 85 | 100 | 100 | 0 | 0 | 0 |
| 0 | 0 | 86 | 100 | 100 | 0 | 0 | 0 |
| 0 | 0 | 87 | 100 | 100 | 0 | 0 | 0 |
| 0 | 0 | 88 | 100 | 100 | 0 | 0 | 0 |
| 0 | 0 | 89 | 100 | 100 | 0 | 0 | 0 |
| 0 | 0 | 90 | 100 | 100 | 0 | 0 | 0 |
| 0 | 0 | 91 | 100 | 100 | 0 | 0 | 0 |
| 0 | 0 | 92 | 100 | 100 | 0 | 0 | 0 |
| 0 | 0 | 93 | 100 | 100 | 0 | 0 | 0 |
| 0 | 0 | 94 | 100 | 100 | 0 | 0 | 0 |
| 0 | 0 | 95 | 100 | 100 | 0 | 0 | 0 |
| 0 | 0 | 96 | 100 | 100 | 0 | 0 | 0 |
| 0 | 0 | 97 | 100 | 100 | 0 | 0 | 0 |
| 0 | 0 | 98 | 100 | 100 | 0 | 0 | 0 |
| 0 | 0 | 99 | 100 | 100 | 0 | 0 | 0 |
| 0 | 0 | 100 | 100 | 100 | 0 | 0 | 0 |

than that met with in other pyranose structures. It is interesting to observe that the C(1)O(1) bond is significantly shorter than the average and therefore the methyl group does not completely eliminate the anomeric shortening observed in the free sugar (Berman, Chu & Jeffrey, 1967). The variations of C(5)O(5), O(5)O(1) and C(1)O(1) from the mean (+0.018, -0.011, -0.025 Å), has the same pattern as that found in methyl α -glucoside (+0.010, -0.10, -0.013 Å), (Berman & Kim, 1968). These discrepancies, and the greater variation of the C-O bond lengths about the mean, as compared with the C-C bond lengths, may be significant and associated with the type of hydrogen bond involved although no correlation is apparent in methyl α -mannoside. A more careful analysis taking account of such factors as extinction, absorption and bonding electrons will be needed to determine to what extent the variations are genuine. These remarks apply also to the CH and OH, bond lengths, the mean values of which are 1.04 and 0.82 Å and therefore somewhat shorter than the usually accepted values of 1.09 and 0.96 Å respectively. A correction for riding motion of the light hydrogen atom on the heavier oxygen and carbon atoms (Busing & Levy, 1964) would probably improve the agreement.

Table 3. Bond lengths and angles in methyl α -D-mannopyranoside

| Bond lengths | | | |
|--------------|---------|---------------|--------|
| C(1)-C(2) | 1.517 Å | C(2)-HC(1) | 1.04 Å |
| C(2)-C(3) | 1.518 | C(2)-HC(2) | 1.16 |
| C(3)-C(4) | 1.524 | C(3)-HC(3) | 0.93 |
| C(4)-C(5) | 1.521 | C(4)-HC(4) | 1.10 |
| C(5)-C(6) | 1.518 | C(5)-HC(5) | 1.09 |
| C(1)-O(1) | 1.407 | C(6)-HC(6') | 1.04 |
| C(2)-O(2) | 1.436 | C(6)-HC(6'') | 1.13 |
| C(3)-O(3) | 1.435 | C(7)-HC(7') | 0.88 |
| C(4)-O(4) | 1.445 | C(7)-HC(7'') | 1.00 |
| C(5)-O(5) | 1.450 | C(7)-HC(7''') | 1.05 |
| C(6)-O(6) | 1.417 | O(2)-HO(2) | 0.67 |
| C(1)-O(5) | 1.421 | O(3)-HO(3) | 0.80 |
| C(7)-O(1) | 1.442 | O(4)-HO(4) | 0.96 |
| | | O(6)-HO(6) | 0.85 |
| Bond angles | | | |
| O(1)C(1)O(5) | 111.7° | C(3)C(4)C(5) | 110.0° |
| C(2)C(1)O(1) | 106.4 | C(3)C(4)O(4) | 109.1 |
| C(2)C(1)O(5) | 110.7 | C(5)C(4)O(4) | 107.2 |
| C(1)C(2)C(3) | 110.7 | C(4)C(5)C(6) | 113.3 |
| C(1)C(2)O(2) | 105.1 | C(4)C(5)O(5) | 110.1 |
| C(3)C(2)O(2) | 111.4 | C(6)C(5)O(5) | 106.0 |
| C(2)C(3)C(4) | 110.8 | C(5)C(6)O(6) | 112.8 |
| C(2)C(3)O(3) | 109.9 | C(1)O(1)C(7) | 113.4 |
| C(4)C(3)O(3) | 107.9 | C(1)O(5)C(5) | 114.3 |

Dihedral angles around the pyranose ring were between 53.5 and 59.3° and therefore in the range found

for other pyranose rings (Kim & Jeffrey, 1967). Also consistent with other carbohydrate structures is the 114.3° angle at the ring oxygen atom, O(5), and the mean value of 110.0° for the carbon valence angles. In the latter there is no significant variation with type, viz. CCO, OCO, and CCC, from the tetrahedral angle of 109.5°. The bond angles involving hydrogen were also normal: an average of 110.0° with a range of 108 to 114°.

The plane of the molecular ring comprising C(1), C(2), C(3), C(4), C(5) and O(5) is defined by the equation:

$$0.0661x - 0.9736y + 0.2185z = 0.4774.$$

Four of the atoms are 0.23₁ Å from this plane; the remaining two, C(1) and C(2), both of which have substituents that are axial, are marginally closer to the ring plane at ± 0.21 , Å - the calculated standard deviations are 0.005 Å.

The oxygen-oxygen contact distances ≤ 3.25 Å are listed in Table 4 together with O...HO distances and O...H-O angles. The hydrogen bonding system comprises two short terminated chains: O(2')H \rightarrow O(1), O(6'')H \rightarrow O(3)H \rightarrow O(4''')H \rightarrow O(5'''''). Two oxygen atoms [O(2) and O(6)] are donors only, two [O(3) and O(4)] are both donors and acceptors, and two [O(1) and O(5)] are acceptors only. One distance [O(3)...O(4)] is characterized by being significantly shorter than the others and at the same time possesses an almost linear arrangement of atoms (174°). The longest oxygen-oxygen contact distance [O(4)...O(5)] on the other hand has the least linear atomic arrangement (151°). All four contacts suggested as hydrogen bonds have H...O distances which are approximately 0.5 Å less than the sum, 2.60 Å, of the van der Waals radii of hydrogen and oxygen (Pauling, 1960). As a group, the oxygen-oxygen contact distances in methyl α -mannoside tend to a longer average distance than in many other carbohydrates (Jeffrey & Rosenstein, 1964).

Although C(7) has a high thermal parameter its associated hydrogen atoms were clearly resolved and indicate that at room temperature the methyl group is not free to rotate. The contact distances of atoms around this group are consistent with van der Waals radii with no unusually close contacts which may help to stabilize the conformation (Mathieson & Poppleton, 1966). The location of the methyl group with respect to the ring is the same as that found in methyl α -glucoside.

The primary hydroxyl group, however, differs in its position. Thus in methyl α -mannoside the C(6)-O(6) bond is approximately perpendicular to the general

Table 4. Hydrogen bond distances (≤ 3.25 Å) and angles in methyl α -D-mannopyranoside

| Distances | | Angles | |
|-------------|---------|------------------|------|
| O(1)...O(2) | 2.872 Å | O(1)...HO(2) | 163° |
| O(3)...O(6) | 2.844 | O(3)...HO(6)O(6) | 156 |
| O(4)...O(3) | 2.776 | O(4)...HO(3)O(3) | 174 |
| O(5)...O(4) | 2.937 | O(5)...HO(4)O(4) | 151 |
| | | O(1)...HO(2)O(2) | 163° |
| | | O(3)...HO(6)O(6) | 156 |
| | | O(4)...HO(3)O(3) | 174 |
| | | O(5)...HO(4)O(4) | 151 |

plane of the molecule and consequently O(6) is *trans* to HC(5) whilst in methyl α -glucoside O(6) is approximately in the plane of the ring and *trans* to C(4) (Berman & Kim, 1968).

We are indebted to Dr A. McL. Mathieson* for helpful discussion and encouragement during the course of this work, and would like also to thank Dr J. C. B. White,† Dr D. A. Wright,‡ and Mr N. Kaye* for providing a number of essential computer programs.

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The Crystal and Molecular Structure of 5-Iodouridine

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The crystal structure of 5-iodouridine ($C_9H_{11}IN_2O_6$) has been determined from data collected on a Hilger and Watts linear diffractometer. The crystals are monoclinic, space group $P2_1$, with cell dimensions: $a = 4.683 \pm 0.005$, $b = 17.14 \pm 0.01$, $c = 14.58 \pm 0.01$ Å and $\beta = 90.85 \pm 0.1^\circ$. The structure was solved by the heavy-atom method and the positional and thermal parameters were refined by the method of least-squares. The final R value for 2143 observed reflexions is 0.061. There are two molecules in the asymmetric unit and the dihedral angle between the base and sugar planes in both is 69° . However, the glycosidic torsion angle, φ_{CN} , is -13.2° in molecule I and -58.7° in molecule II. In molecule I atom C(3') of the sugar ring is displaced 0.54 Å *endo*, and in molecule II atom C(2') is displaced 0.63 Å *endo*. The orientation of the C(5')-O(5') bond is *gauche* to both the C(4')-O(1') and C(4')-C(3') bonds in molecule I, but in molecule II it is *trans* to C(4')-O(1') and *gauche* to C(4')-C(3').

Introduction

The structure determination of 5-iodouridine (IUR) (Fig. 1) was undertaken as part of a series of structure determinations of nucleic acid components and their analogues. Single-crystal studies of nucleosides and nucleotides give important information about the bond lengths and angles, and about the conformations of the molecules, which can be used when molecular models of polynucleotide structures are constructed. It was mainly to obtain information about the conformation of the molecule that IUR was studied. Knowledge of the crystal and molecular structure of IUR was also desirable as a first step in the study of the effects of ionizing radiations on single crystals of IUR by electron-spin resonance. Such studies have already been made on 5-iodo-2'-deoxyuridine (Nice &

Rorke, 1969). It is hoped that similar studies will eventually elucidate the effects of ionizing radiations on the nucleic acids themselves, and explain the change in sensitivity when certain analogues are incorporated into the structure in place of a normal component. A preliminary account of the conformations of the molecules has been given recently (Rahman & Wilson, 1970).

Experimental

Crystal of 5-iodouridine ($C_9H_{11}IN_2O_6$) were obtained by evaporation from aqueous solutions. The unit-cell dimensions were first obtained from Weissenberg photographs and were then refined on a Wooster four-circle diffractometer using Cu $K\alpha$ radiation. The crystals are monoclinic and elongated along the a axis. The unit-cell dimensions are: