

## The Crystal Structure of 1,6-Anhydro- $\beta$ -D-glucopyranose

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The crystal structure of 1,6-anhydro- $\beta$ -D-glucopyranose,  $C_6H_{10}O_5$ , levoglucosan, has been determined by the direct method using the tangent formula for phase determination. The structure was refined by anisotropic least-squares using Cu  $K\alpha$  data to give a final  $R$  value of 0.035 for 687 reflections. The space group is  $P2_12_12_1$ , with  $Z=4$  and unit-cell dimensions  $a=6.684(2)$ ,  $b=13.266(7)$ ,  $c=7.547(2)$  Å. The molecule has a dicyclic structure, with an anhydro bridge across a pyranose ring forming a boat-shaped seven-membered ring. The three hydroxyl substituents are axial relative to the pyranose ring, two *exo*-axial at C(2) and (4) and one *endo*-axial at C(3). The pyranose ring has the  $1C$  chair conformation which is distorted by the formation of the anhydride, so as to compress C(1) and C(5) and extend C(2) and C(4). The pyranose ring oxygen angle is less and the two *syn*-hydroxyl groups at C(2) and C(4) are further apart than in an unstrained pyranose ring system.

### Introduction

1,6-Anhydro- $\beta$ -D-glucopyranose, levoglucosan,  $C_6H_{10}O_5$  is an internal glycoside with a dicyclic ring system, which is formed by the dry distillation of D-glucose. It is one of the more important anhydro sugars (Peat, 1946), since it is easily hydrolyzed to glucose by aqueous acid, and can therefore be used in the synthesis of glucose derivatives substituted in the positions 2, 3 or 4 and in the polymerization to dextrans (Carvalho, Prins & Schuerch, 1959).

The anhydride formation must take place through conversion of D-glucose to its less stable  $1C$  conformation, as shown in Fig. 1. The anhydro molecule *IIa* has three hydroxyls in the axial orientation, and it is not immediately obvious that this conformer will be preferred over that shown in *IIb*, in which the pyranose ring is boat-shaped and the hydroxyl groups are equatorially oriented.

The structure was determined in order to establish the conformation in the crystalline state and to examine the strain effect of anhydride formation on the shape of the pyranose ring and the orientation of the three hydroxyl groups.

### Experimental

Large, transparent crystals of 1,6-anhydro- $\beta$ -D-glucopyranose were provided by the Chemical Institute of the Slovak Academy of Sciences, Bratislava, Czechoslovakia. The crystal data are as follows:

1,6-anhydro- $\beta$ -D-glucopyranose,  $C_6H_{10}O_5$ ,  
M.W. 162.08, m.p. 184°C.

Space group  $P2_12_12_1$ , from systematic absences,  
 $h00$ ,  $h=2n+1$ ;  $0k0$ ,  $k=2n+1$ ,  $00l$ ,  $l=2n+1$ .

$$a = 6.684(2) \text{ \AA}$$

$$b = 13.266(7)$$

$$c = 7.547(2)$$

$$Z = 4$$

$$D_m = 1.618 \text{ g.cm}^{-3}, D_x = 1.608 \text{ g.cm}^{-3}$$

$$\mu_{Cu K\alpha} = 12.45 \text{ cm}^{-1}$$

The cell parameters and three-dimensional intensity data were measured on a Picker FACS I diffractometer using Cu  $K\alpha$  radiation with a crystal of dimensions  $0.3 \times 0.3 \times 0.25$  mm mounted along the diagonal axis [101]. A  $2\theta$  scanning mode with varying interval was used up to  $2\theta = 130^\circ$ . Of the 687 recorded data,

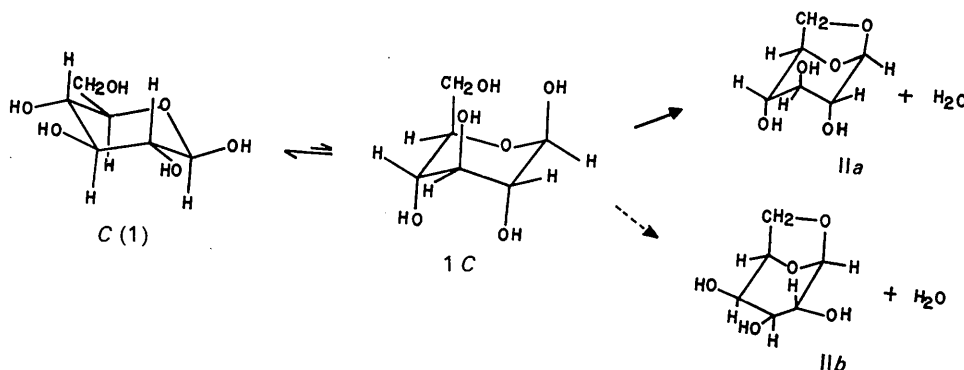


Fig. 1. Formation of 1,6-anhydro- $\beta$ -D-glucopyranose from  $\beta$ -D-glucose.

Table 1. Fractional atomic coordinates and anisotropic thermal parameters in 1,6-anhydro- $\beta$ -D-glucopyranose

Key to the atomic numbering is given in Fig. 2. The temperature factor expression used was  $\exp \{-(h^2\beta_{11} + k^2\beta_{22} + l^2\beta_{33} + 2hk\beta_{12} + 2hl\beta_{13} + 2kl\beta_{23})\}$ .

The estimated standard deviations are in parentheses.

	x	y	z	$\beta_{11}$	$\beta_{22}$	$\beta_{33}$	$\beta_{12}$	$\beta_{13}$	$\beta_{23}$
C(1)	0.1984 (5)	0.0482 (2)	0.3060 (5)	0.0160 (6)	0.0024 (1)	0.0138 (6)	0.0009 (3)	-0.0001 (6)	0.0009 (3)
C(2)	0.2536 (4)	0.1311 (2)	0.1761 (4)	0.0101 (6)	0.0029 (1)	0.0104 (5)	0.0000 (2)	0.0009 (5)	-0.0003 (2)
C(3)	0.0717 (4)	0.1964 (2)	0.1275 (4)	0.0124 (6)	0.0025 (1)	0.0089 (4)	-0.0003 (3)	-0.0012 (5)	0.0005 (2)
C(4)	-0.0730 (4)	0.2105 (2)	0.2844 (4)	0.0100 (6)	0.0026 (1)	0.0120 (5)	-0.0000 (2)	-0.0006 (5)	0.0000 (2)
C(5)	-0.0834 (5)	0.1170 (2)	0.3995 (4)	0.0136 (7)	0.0029 (2)	0.0128 (5)	-0.0006 (3)	0.0036 (5)	0.0002 (2)
C(6)	-0.1427 (5)	0.0231 (2)	0.2947 (6)	0.0162 (8)	0.0028 (2)	0.0171 (7)	-0.0018 (3)	0.0032 (7)	-0.0001 (3)
O(1)	0.0472 (4)	0.0153 (2)	0.2330 (3)	0.0191 (6)	0.0022 (1)	0.0182 (5)	-0.0004 (2)	0.0033 (5)	-0.0013 (2)
O(2)	0.4133 (3)	0.1900 (2)	0.2469 (3)	0.0121 (5)	0.0043 (1)	0.0102 (4)	-0.0015 (2)	-0.0001 (3)	0.0003 (2)
O(3)	-0.0278 (4)	0.1497 (2)	0.0180 (3)	0.0192 (5)	0.0038 (1)	0.0109 (4)	-0.0010 (2)	-0.0049 (4)	0.0002 (2)
O(4)	-0.0055 (4)	0.2903 (1)	-0.3966 (3)	0.0186 (5)	0.0023 (1)	0.0115 (4)	-0.0005 (2)	0.0015 (4)	-0.0004 (2)
O(5)	0.1149 (3)	0.0903 (2)	0.4590 (3)	0.0172 (5)	0.0032 (1)	0.0103 (3)	0.0002 (2)	-0.0009 (4)	0.0014 (2)
H(1)	0.333	0.003	0.339	0.062					
H(2)	0.305	0.109	0.062	0.086					
H(3)	0.124	0.266	0.225	0.225					
H(4)	-0.201	0.220	0.506	0.506					
H(5)	-0.170	0.126	0.384	0.384					
H(6)	-0.192	-0.024	0.192	0.192					
H(O3)	0.227	0.045	0.369	0.369					
H(O2)	0.407	0.205	0.097	0.097					
H(O3)	-0.061	0.200	-0.352	-0.352					
H(O4)	-0.035	0.349							

643 reflections were observed above background. The data were reduced to structure amplitudes using an IBM 1130 program (Shiono, 1969). No absorption corrections were applied. The crystal density was measured by the flotation method in a mixture of carbon tetrachloride and bromoform at 23°C.

Determination and refinement of the structure

The structure was solved by the application of the tangent formula (Karle & Hauptman, 1956) using an IBM 7090 version of the Hall (1968) direct phasing methods. Four reflections were selected as starting phases, three to define the origin (3, 11, 0; 5, 8, 0; 0, 2, 1) and one to define the enantiomorph (0, 15, 2) with E's of 2.90, 2.25, 1.82 and 1.94 respectively. The tangent formula refinement was then applied to 148 reflections with E value > 1.40. The eleven highest peaks on the resulting E map corresponded to the eleven oxygen and carbon atoms of the molecule. The first trial structure gave an R value of 0.30 for all reflections. Refinement was by several cycles of isotropic and of anisotropic structure factor calculations using the block-diagonal least-squares program for IBM 1130 (Shiono, 1968) and the full-matrix least-squares program for IBM 7090 (Shiono, 1966). The function minimized was  $\omega_i(K|F_o| - |F_c|)^2$  where  $\omega_i = |F_o|^{-2}$  for  $|F_o| \geq 4|F_{min}|$  and  $\omega_i = 4|F_{min}|^{-2}$  for  $|F_o| < 4|F_{min}|$ , with  $|F_{min}| = 8.4$ . The final cycle gave an R value of 0.035 and an average  $\Delta/\sigma$  of 0.7. All the hydrogen atoms were clearly located on a difference synthesis at an R value of 0.090. Their coordinates were not refined and they were assigned the thermal parameters of the atoms to which they are attached. Atomic parameters are given in Table 1. The observed and calculated structure factors are

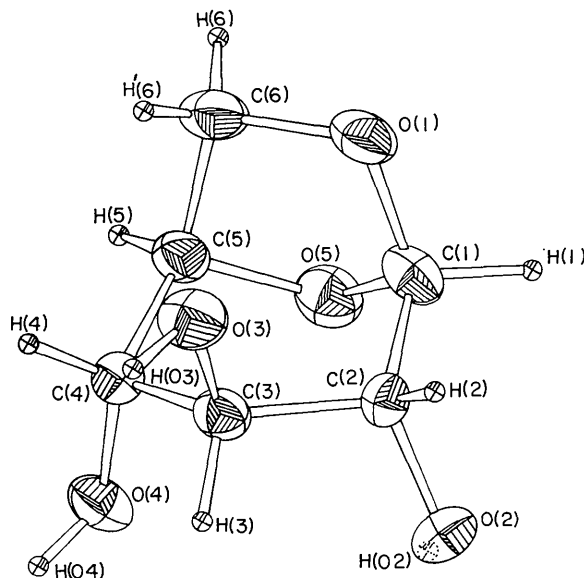


Fig. 2. Molecular conformation and atomic numbering of 1,6-anhydro- $\beta$ -D-glucopyranose. ORTEP plot of thermal ellipsoids.

Table 2. Observed and calculated structure factors

Columns are: Index, 10|F<sub>obs</sub>|, 10|F<sub>calc</sub>|, 10A<sub>calc</sub>, 10B<sub>calc</sub>.

\* Indicates unobserved reflections.

Index	10 F <sub>obs</sub>	10 F <sub>calc</sub>	10A <sub>calc</sub>	10B <sub>calc</sub>
2 430 452 452-0	0	0	0	0
2 250 250 250-0	0	0	0	0
8 92 95 95-0	0	0	0	0
8 65 63 63-0	0	0	0	0
1 96 97 0 97-	0	0	0	0
2 220 223 0 223-	0	0	0	0
3 95 103 0 103-	0	0	0	0
4 110 118 0 118-	0	0	0	0
5 110 114 0 114-	0	0	0	0
6 140 141 0 141-	0	0	0	0
8 18 19 0 19-	0	0	0	0
7 127 127 0 127	0	0	0	0
1 146 149 149-0	0	0	0	0
1 695 739 739-0	0	0	0	0
2 184 190 190-0	0	0	0	0
3 93 92 0 92-	0	0	0	0
4 166 174 174-0	0	0	0	0
5 44 41 0 41-	0	0	0	0
6 72 73 73-0	0	0	0	0
8 110 106 106-0	0	0	0	0
7 16 15 15-0	0	0	0	0
1 207 213 0 213-	0	0	0	0
2 294 300 0 300-	0	0	0	0
3 53 53 0 53-	0	0	0	0
4 164 158 0 158-	0	0	0	0
5 16 14 0 14-	0	0	0	0
6 102 103 0 103-	0	0	0	0
8 14 14 0 14-	0	0	0	0
7 22 20 0 20-	0	0	0	0
0 -77 68 37 0	0	0	0	0
1 148 141 141-0	0	0	0	0
2 632 637 637-0	0	0	0	0
3 437 428 428-0	0	0	0	0
5 6 5 0 5-	0	0	0	0
6 131 133 133-0	0	0	0	0
8 21 20 0 20-	0	0	0	0
7 128 132 132-0	0	0	0	0
1 351 361 0 361-	0	0	0	0
2 308 303 0 303-	0	0	0	0
3 60 63 63-0	0	0	0	0
4 264 264 0 264-	0	0	0	0
5 225 226 0 226-	0	0	0	0
8 9 9 0 9-	0	0	0	0
7 27 26 0 26-	0	0	0	0
0 321 323 323-0	0	0	0	0
1 150 154 154-0	0	0	0	0
2 313 300 300-0	0	0	0	0
3 341 328 328-0	0	0	0	0
4 146 146 146-0	0	0	0	0
5 51 54 54-0	0	0	0	0
6 8 6 6-0	0	0	0	0
8 7 7 7 0 7-	0	0	0	0
7 37 37 0 37-	0	0	0	0
1 163 165 0 165-	0	0	0	0
2 77 71 0 71-	0	0	0	0
3 197 198 0 198-	0	0	0	0
4 164 162 0 162-	0	0	0	0
5 121 124 124-0	0	0	0	0
6 150 153 0 153-	0	0	0	0
7 64 65 0 65-	0	0	0	0
0 3 3 0 3-	0	0	0	0
1 68 67 67-0	0	0	0	0
2 75 66 66-0	0	0	0	0
4 20 18 18-0	0	0	0	0
5 160 153 153-0	0	0	0	0
6 61 58 58-0	0	0	0	0
7 28 28 0 28-	0	0	0	0
1 170 171 0 171-	0	0	0	0
2 5 5 0 5-	0	0	0	0
3 118 147 0 147-	0	0	0	0
4 40 42 0 42-	0	0	0	0
5 29 29 0 29-	0	0	0	0
7 19 14 0 14-	0	0	0	0
0 261 268 268-0	0	0	0	0
1 6 6 0 6-	0	0	0	0
2 81 88 88-0	0	0	0	0
3 30 24 24-0	0	0	0	0
4 37 39 39-0	0	0	0	0
5 88 92 92-0	0	0	0	0
6 146 139 139-0	0	0	0	0
1 138 142 0 142-	0	0	0	0
2 6 2 0 2-	0	0	0	0
4 69 70 0 70-	0	0	0	0
5 38 37 0 37-	0	0	0	0
6 5 5 0 5-	0	0	0	0
0 196 203 203-0	0	0	0	0
1 46 45 45-0	0	0	0	0
2 116 120 120-0	0	0	0	0
3 89 84 84-0	0	0	0	0
4 103 105 105-0	0	0	0	0
5 43 38 38-0	0	0	0	0
1 29 29 0 29-	0	0	0	0
2 6 14 0 14-	0	0	0	0
3 25 29 0 29-	0	0	0	0
4 47 50 0 50-	0	0	0	0
1 72 72 0 72-	0	0	0	0
1 147 151 151-0	0	0	0	0
2 20 24 24-0	0	0	0	0
3 67 69 69-0	0	0	0	0
1 29 19 0 19-	0	0	0	0
2 102 100 0 100-	0	0	0	0
1 696 801 801-0	0	0	0	0
2 238 238 238-0	0	0	0	0
3 231 229 0 229-	0	0	0	0
4 126 130 130-0	0	0	0	0
5 35 29 0 29-	0	0	0	0
6 49 48 48-0	0	0	0	0
8 44 45 45-0	0	0	0	0
7 70 72 0 72-	0	0	0	0
1 335 344 0 344-	0	0	0	0

Table 3. *Least-squares planes in 1,6-anhydro-β-D-glucopyranose*Equation for plane:  $Ax + By + Cz = D$ , where  $x, y, z$  are in Å.

	Atoms in plane	Distance from best plane	Constants	
Pyranose ring	C(1)	0.001	A	0.501
	C(2)	-0.001	B	0.477
	C(4)	0.001	C	0.722
	C(5)	-0.001	D	2.638
		C(3)	-0.460	
Anhydride ring		O(5)	0.820	
	C(1)	0.012	A	0.049
	C(5)	-0.011	B	-0.563
	C(6)	0.017	C	0.825
	O(1)	-0.018	D	1.599
		O(5)	0.623	
	C(2)	0.019	A	0.019
	O(2)	-0.015	B	-0.610
	C(4)	-0.018	C	0.792
	O(4)	0.014	D	0.005

Table 4. *Values of A, B, B' and μ used in the energy function*Energies are expressed in kcal.mole<sup>-1</sup>.

	A	B (× 10 <sup>-4</sup> )	B' (× 10 <sup>-4</sup> )	μ
H ··· H	49	0.4682	1.0000	4.60
H ··· O <sub>h</sub> *	93	1.4365	0.7978	4.12
H ··· O <sub>r</sub> *	100	1.5446	0.8478	4.12
H ··· C	125	3.0118	1.2900	4.12
O <sub>h</sub> ··· O <sub>h</sub>	201	4.8430	0.6364	3.63
O <sub>h</sub> ··· O <sub>r</sub>	214	5.1562	0.6763	3.63
O <sub>h</sub> ··· C	250	9.1125	1.0278	3.63
O <sub>r</sub> ··· O <sub>r</sub>	227	5.4695	0.7187	3.63
O <sub>r</sub> ··· C	266	9.6957	1.0923	3.63
C ··· C	325	17.4483	1.6600	3.63

\* O<sub>h</sub> and O<sub>r</sub> represent hydroxyl and ring oxygen, respectively.Table 5. *Bond lengths and angles in 1,6-anhydro-β-D-glucopyranose*

The estimated standard deviations given in parentheses refer to the last decimal positions at respective values.

i	j	D(ij)	i	j	k	∠(ijk)
C(1)	C(2)	1.518 (4) Å	C(1)	C(2)	C(3)	111.7 (2)°
C(2)	C(3)	1.537 (4)	C(2)	C(3)	C(4)	112.4 (2)
C(3)	C(4)	1.540 (4)	C(3)	C(4)	C(5)	111.7 (2)
C(4)	C(5)	1.516 (5)	C(4)	C(5)	C(6)	112.5 (3)
C(5)	C(6)	1.527 (5)	C(4)	C(5)	O(5)	109.6 (3)
			C(5)	O(5)	C(1)	102.0 (2)
C(2)	O(2)	1.426 (3)	O(5)	C(1)	C(2)	110.0 (2)
C(3)	O(3)	1.425 (4)	O(1)	C(1)	C(2)	110.5 (2)
C(4)	O(4)	1.429 (4)	C(5)	C(6)	O(1)	103.1 (3)
			C(6)	O(1)	C(1)	106.8 (2)
C(1)	O(1)	1.427 (4)	O(1)	C(1)	O(5)	105.8 (2)
C(6)	O(1)	1.444 (4)	O(5)	C(5)	C(6)	101.5 (3)
C(1)	O(5)	1.399 (4)				
C(5)	O(5)	1.443 (4)	C(1)	C(2)	O(2)	109.6 (2)
			O(2)	C(2)	C(3)	111.9 (2)
Mean value			C(2)	C(3)	O(3)	107.9 (2)
C-C	1.528 (11)		O(3)	C(3)	C(4)	110.6 (2)
C-O(H)	1.427 (2)		C(3)	C(4)	O(4)	110.3 (2)
C-O	1.428 (21)		O(4)	C(4)	C(5)	106.4 (2)
			Mean value			
			C-C-C			112.1 (4)
			C-C-O(H)			109.5 (18)
			Five-membered ring formed by			
			C(1)-O(1)-C(6)-C(5)-O(5)			103.8 (24)

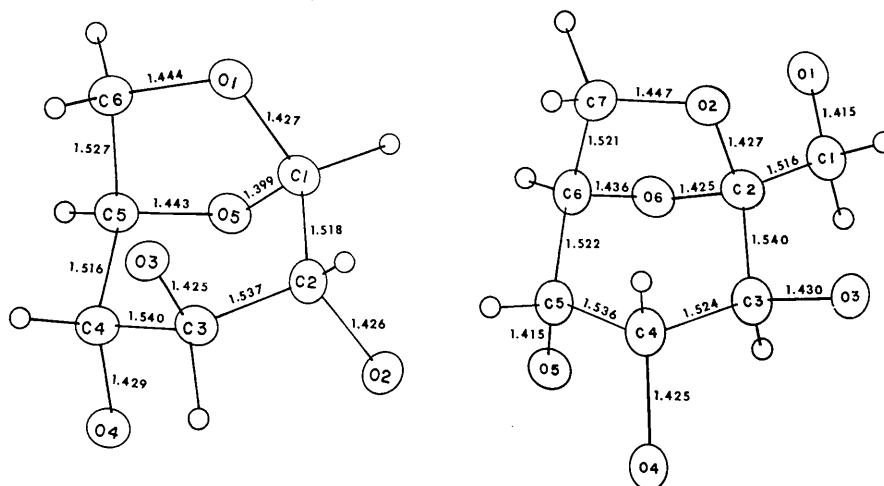


Fig. 3. Comparison of bond lengths in 1,6-anhydro- $\beta$ -D-glucopyranose (left) and in sedoheptulosan (right) (Brown & Thiessen, 1969).

given in Table 2. The atomic scattering factors used were those of Cromer & Waber (1965) for carbon and oxygen and of Stewart, Davidson & Simpson (1965) for hydrogen.

### Description of the structure

The molecular conformation and atomic numbering are shown in Fig. 2. The molecule has the conformation shown in II*a*, in which the pyranose ring has the 1C chair form with the hydroxyl groups axial at C(2), C(3) and C(4). The seven-membered ring which includes the six carbon atoms and O(1) is boat-shaped with the atoms lying in the three planes containing the atoms C(1) C(5) C(6) O(1), C(1) C(2) C(4) C(5), (see Table 3) and C(2) C(3) C(4), which make dihedral angles about C(1)···C(5) and C(2)···C(4) of 110.7° and 147.4°, respectively. The atoms C(2), O(2), C(4), O(4) are also in the same plane within 0.019 Å. In the unstrained conformation, the bonds C(2)–O(2) and C(4)–O(4), C(3)–O(3) and C(5)–C(6) will be eclipsed and parallel, thereby bringing O(2) and O(4), O(3) and C(6) to within non-bonded distances of 2.5 Å. These distances correspond to non-bonding interactions which in carbohydrates normally lead to instability relative to some alternate conformation, in this case the conformer shown in II*b*. However, in this molecule, the pyranose ring is strained by the formation of the anhydro ring in a manner such that the separations of O(2) and O(4), O(3) and C(6) are increased to 3.30 and 3.00 Å, respectively. The greater stability due to this reduction of the non-bonding repulsion by anhydro ring formation has been referred to as the 'reflex effect' by Sandris & Ourisson (1958).

The alternate conformation, II*b*, in which the pyranose has the boat form, the seven-membered ring is chair-shaped, and the substituent hydroxyl groups are all equatorial, might still be competitive in energy. A

calculation was therefore made of the non-bonding interaction energies for all atom pairs separated by more than two covalent bonds and closer than 4 Å,

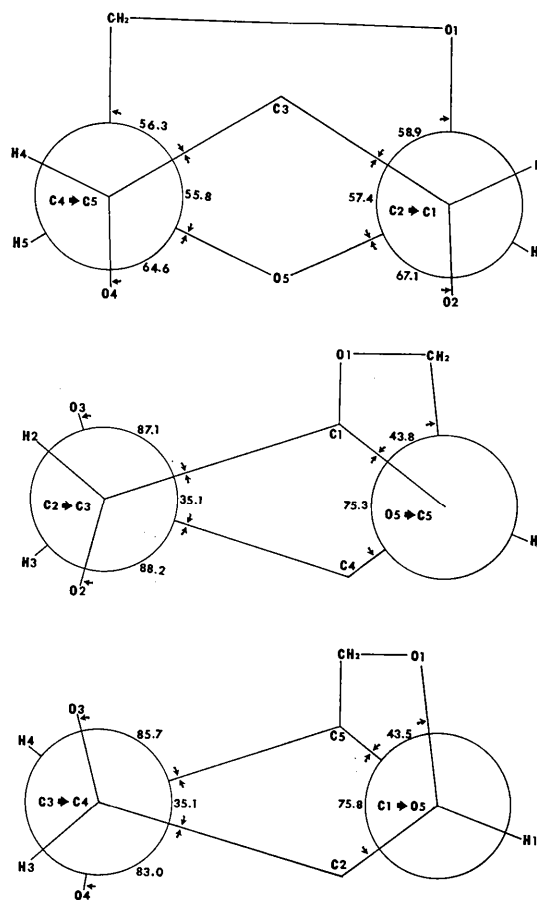


Fig. 4. The conformation angles about the C-C and C-O bonds of the pyranose ring in 1,6-anhydro- $\beta$ -D-glucopyranose.

Table 6. *Hydrogen bonding distances and angles*

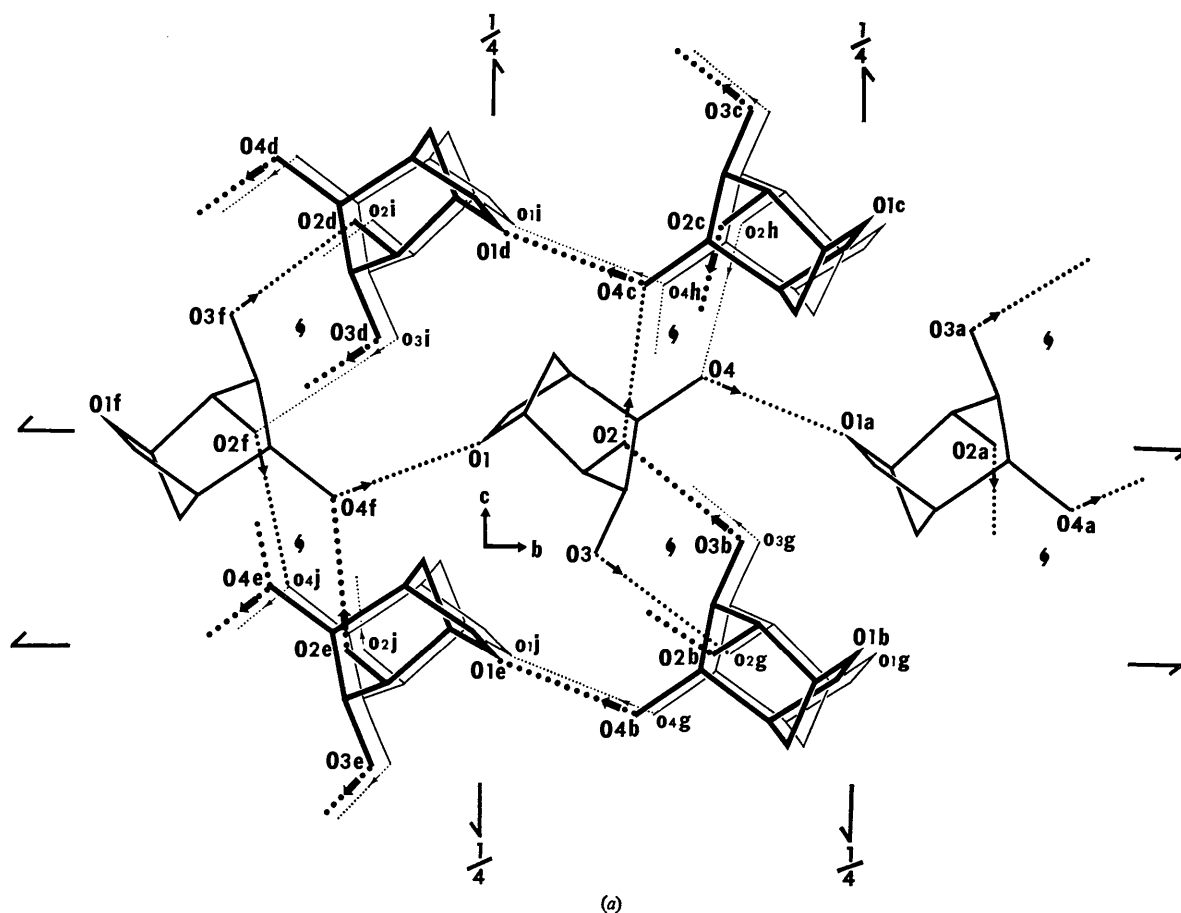
<i>i</i>	<i>j</i>	<i>k</i>	<i>D</i> ( <i>ik</i> )	$\angle$ ( <i>ijk</i> )	<i>i</i>	<i>j</i>	<i>k</i>	<i>D</i> ( <i>jk</i> )	$\angle$ ( <i>ijk</i> )
O(2)	H(O2)	O(4c)	2.756 Å	157°	C(2)	O(2)	O(4c)	2.756 Å	124.4°
O(3)	H(O3)	O(2g)	2.768	170	C(3)	O(3)	O(2g)	2.768	102.3
O(4)	H(O4)	O(1a)	2.771	168	C(4)	O(4)	O(1a)	2.771	116.7
O(2)	H(O2)	O(5)	2.878	100	C(2)	O(2)	O(5)	2.878	55.8

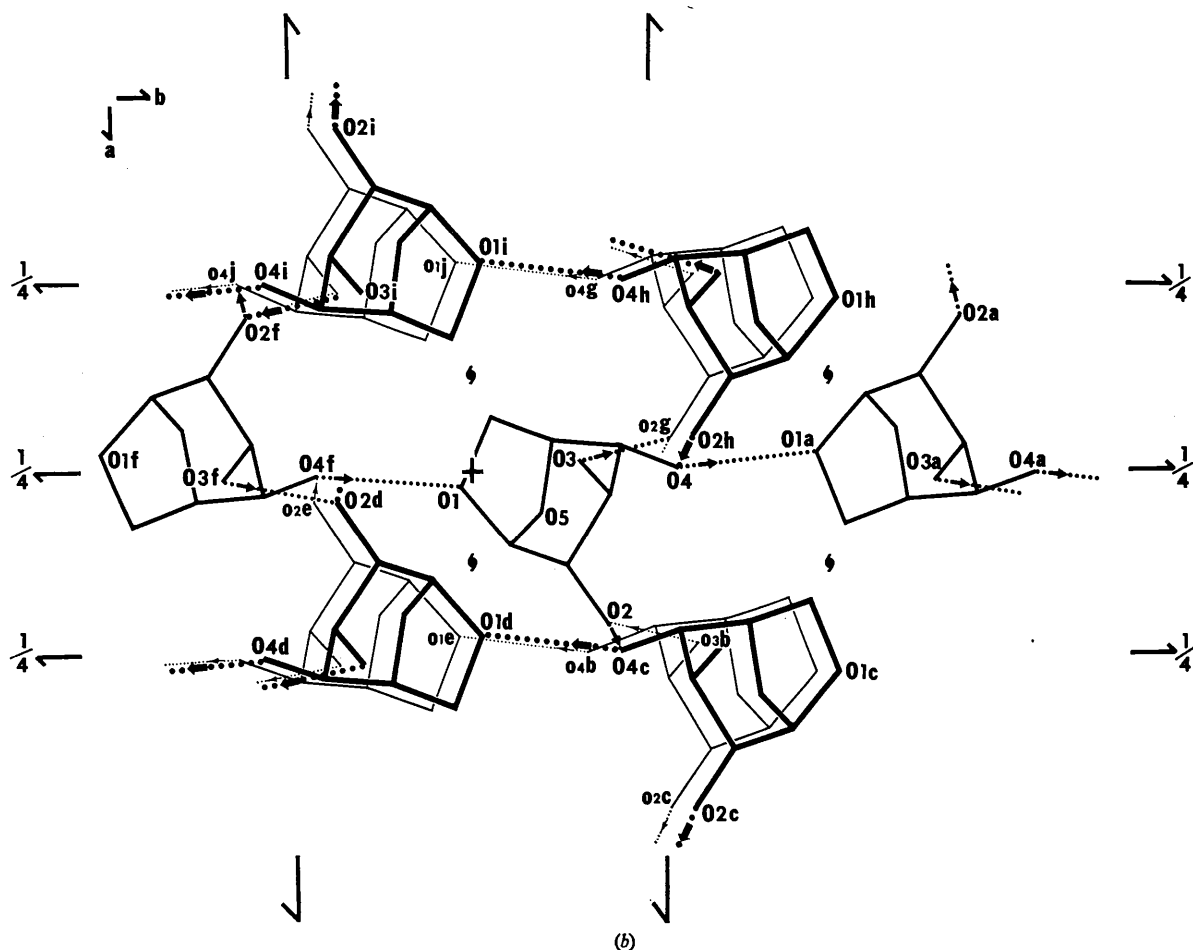
Non-bonded O...O distances less than 3.3 Å

O(1)	O(5e)	3.210 Å
O(2)	O(4)	3.299
O(3)	O(1)	2.938
O(4)	O(5)	2.812
O(4)	O(5h)	3.182

Symmetry code

No symbol	<i>x</i>	<i>y</i>	<i>z</i>
<i>a</i>	- <i>x</i>	$\frac{1}{2}+y$	$\frac{1}{2}-z$
<i>b</i>	$\frac{1}{2}+x$	$\frac{1}{2}-y$	- <i>z</i>
<i>c</i>	$\frac{1}{2}+x$	$\frac{1}{2}-y$	1- <i>z</i>
<i>d</i>	$\frac{1}{2}-x$	- <i>y</i>	$\frac{1}{2}+z$
<i>e</i>	$\frac{1}{2}-x$	- <i>y</i>	$-\frac{1}{2}+z$
<i>f</i>	- <i>x</i>	$-\frac{1}{2}+y$	$\frac{1}{2}-z$
<i>g</i>	$-\frac{1}{2}+x$	$\frac{1}{2}-y$	- <i>z</i>
<i>h</i>	$-\frac{1}{2}+x$	$\frac{1}{2}-y$	1- <i>z</i>
<i>i</i>	$-\frac{1}{2}-x$	- <i>y</i>	$\frac{1}{2}+z$
<i>i</i>	$-\frac{1}{2}-x$	- <i>y</i>	$-\frac{1}{2}+z$

Fig. 5. The hydrogen bonding in the crystal structure of 1,6-anhydro- $\beta$ -D-glucopyranose. Solid lines indicate the molecules; dotted lines indicate the hydrogen bonds with the arrows pointing in the donor direction. (a) View down the *a* axis.

Fig 5 (cont.) (b) View down the *c* axis.

by means of the two commonly used Lennard-Jones type functions;

$$U_{ij}(r_{ij}) = -A/r_{ij}^6 + B/r_{ij}^{12}$$

and

$$U_{ij}(r_{ij}) = -A/r_{ij}^6 + B' \exp \{-\mu r_{ij}\}.$$

The values for  $A$ ,  $B$ ,  $B'$  and  $\mu$  were calculated according to the methods suggested by Pitzer & Catalano, (1956) and Hendrickson (1961), with van der Waals radii for carbon, oxygen and hydrogen of 1.6, 1.4 and 1.25 Å respectively. The values of these constants are given in Table 4. Both expressions gave non-bonding interaction energies favoring the observed axially substituted conformer by 1.1 and 0.7 kcal.mole<sup>-1</sup> respectively. The largest repulsion interaction in the conformer IIb was between the ring oxygen and C(3), *i.e.*, that characteristic of the boat-shaped pyranose ring.

The intramolecular distances and valence angles are given in Table 5. The average values of C-C and C-O(H) bond lengths and C-C-C and C-C-O(H) angles are 1.528 (11), 1.427 (2) Å and 112.1 (4), 109.5

(18)° respectively. These values are not significantly different from those observed in other pyranose sugars (Berman, Chu & Jeffrey, 1967). The same ring system has been studied by Brown & Thiessen (1969) in sedoheptulosan in the crystal structure of the monohydrate. The bond distances are compared in Fig. 3.

It is most interesting to note that these bond distances agree to better than 0.015 Å, except for those which include C(1) where the configuration of the two molecules is different. There we observe differences of 0.026 Å in C(1)-O(5) and 0.022 Å in C(1)-C(2), both being longer in the molecule which has the primary alcohol group on C(1). This observation is of marginal significance and requires additional evidence before it can be regarded as a characteristic of the anhydro-pyranose carbohydrate derivatives.

The chair conformation of the pyranose ring in this anhydro sugar is significantly strained. The conformation angles, shown in Fig. 4, differ by as much as 25° from those observed in unstrained pyranose rings (Berman & Kim, 1968). While the dihedral angles around C(1)-C(2) and C(4)-C(5) are close to the unstrained values, those around C(2)-C(3) and C(3)-C(4)

are smaller, and those around C(5)–O(5) and O(5)–C(1) are larger, *i.e.*, 35.1, 35.1 and 75.3, 75.8° respectively. There are corresponding differences in the deviations of C(3) and O(5) from the plane of C(1), C(2), C(4), C(5) (Table 5). The angle at the pyranose ring oxygen, O(5), is about 10° smaller than normal and comparable with that at the oxygen in the anhydro ring.

The hydrogen bonding is shown in Table 6 and Fig. 5(a) and (b). The hydroxyls at O(2) and O(4) form two hydrogen bonds, one donor and one acceptor; that at O(3) forms only a donor bond to O(2). The molecules are linked by finite chains of hydrogen bonds through O(3b)H–O(2)H–O(4c)H–O(1d). The hydrogen H(O2) is 'anti' to H(2) with respect to the C(2)–O(2) bond and O(2)–H(O2) is directed toward the mid-point of the triangle formed by O(5) and O(4) of the same molecule and O(4c) of an adjacent molecule. The distances from H(O2) to O(4c), O(5) and O(4) are 1.86, 2.57 and 2.98 Å. Therefore, only the intermolecular interaction can be regarded as a hydrogen bond. However, there is nuclear magnetic resonance and infrared spectral evidence that this 'anti' conformation of H(O2) occurs in a 5% CCl<sub>4</sub> solution of the 3,4-dimethyl ether derivative and that this may be a characteristic 'conformational' property of certain levoglucosan derivatives (Seib, 1970). This would imply some intramolecular stabilization of the conformation observed in this structure.

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## The Crystal Structure of the $\nu$ Phase, Mn<sub>81.5</sub>Si<sub>18.5</sub>\*

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The crystal structure of the  $\nu$  phase, Mn<sub>81.5</sub>Si<sub>18.5</sub> has been determined by single-crystal X-ray diffraction analysis. The lattice constants are  $a_0 = 16.992$  (4),  $b_0 = 28.634$  (7),  $c_0 = 4.656$  (1) Å, and the space group is *Immm* ( $D_{2h}^{25}$ ), with 186 atoms in the unit cell. The structure has been refined by full-matrix least-squares analysis to a final *R* index of 0.041, based on the 2305 reflections observed with the diffractometer. The  $\nu$  phase is related to the *P* phase, and belongs to the family of  $\sigma$ -phase related, layered, tetrahedrally close-packed structures, with percentages of sites with CN 12, 14, 15 and 16 respectively: 40, 43, 11 and 6. All positions with CN > 12 are occupied by Mn, except two CN 14 positions which have some Si present. The CN 12 positions are occupied by various mixtures of Mn and Si, except three positions which are occupied by Mn only, and two positions which are occupied by Si only. The structure may also be described as a stacking of two kinds of 'tiles' each consisting of 7 hexagons and 12 triangles, with the space in-between filled up by hexagons, pentagons and triangles.

### Introduction

The  $\nu$  phase was described by Gupta (1964) in the Mn–Fe–Si system as a binary Mn–Si phase about one

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at. % wide, centered at 81.5 at. % Mn and extending into the ternary system only up to 1.5 at. % Fe. Earlier the occurrence of a complex phase in the Mn–Si system in this region was mentioned by Åmark, Borén & Westgren (1936) and subsequently called the *N* phase by Kuzma & Hladyshevskii (1964). A specimen of the  $\nu$  phase of stated composition Mn<sub>81.5</sub>Si<sub>18.5</sub>, annealed for 72 hours at 1000°C, was kindly made available to us