The Crystal Structure of Calcium-α-D-glucoisosaccharate and Some Extended Hückel Calculations on α-D-Glucoisosaccharinic Acid

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The crystal structure of Ca- α -D-glucoisosaccharate, Ca(C₆H₁₁O₆)₂, has been refined from three-dimensional X-ray diffraction data by least-squares methods. The crystals are orthorhombic, space group $P2_12_12$, with a=19.609, b=6.782, and c=5.782 Å. The final R index is 0.108. The compound is isostructural with the corresponding strontium salt,¹ which was determined with higher precision (R=0.058). The structure of the α -D-glucoisosaccharate group is not significantly different from that found in the strontium compound and the proposed hydrogen bonds are essentially the same.

Some semiempirical conformational analyses were made on the α-D-glucoisosaccharinic acid using extended Hückel calculations.²⁻⁵ Population analysis and net charges were calculated for the free acid

In connection with a study on α -D-glucoisosaccharinic acid, X-ray data were collected for the isomorphous strontium and calcium glucoisosaccharates with different techniques.

In both cases the data were collected with a General Electric Single-Crystal Orienter equipped with a scintillation detector and a pulse-height analyser set to collect about 90 % of the $\text{Cu}K\alpha$ radiation used. The measuring technique used for the strontium compound was the θ -2 θ scanning technique. It was believed that the peak intensity measurement technique would speed up the data collection. The calcium compound was investigated by this method. In this case, however, it was found necessary to search in 2θ for the maximum intensity positions and no measuring time was saved.

A powder specimen of Ca- α -D-glucoisosaccharate ⁶ was kindly supplied by A. Ishizu (Swedish Forest Products Research Laboratory). Single crystals suitable for X-ray analysis were obtained by slow cooling of a saturated aqueous solution of the salt. The dimensions of the crystal used were $0.01 \times 0.02 \times 0.06$ mm³ with the long edge parallel to the c axis. Approxi-

mately 91 % of the reflections not systematically absent had observable intensities so that 478 reflections with 2θ less than 102° could be measured. The background was measured 20 seconds at \pm 1° from the 2θ value found for the maximum intensity. The peak intensities were measured 40 seconds. For obtaining integrated intensities a calibration curve was determined. It was based on 15 reflections with increasing 2θ values measured with the $\theta-2\theta$ scanning technique. No corrections of the peak intensities for obtaining integrated intensities were found necessary. Lp and absorption corrections ($\mu=39~{\rm cm}^{-1}$) were applied to the net intensity counts.

Powder photographs were taken in a Guinier focusing camera using $CuK\alpha_1$ radiation ($\lambda = 1.54056$ Å) and potassium chloride (a = 6.2930 Å) as an internal standard. The unit-cell dimensions refined by least-squares from the X-ray powder pattern are given in Table 1.

Table 1. Crystallographic data of Ca-a-D-glucoisosaccharate

Lattice constants.	$egin{array}{lll} a &= 19.609 \pm 2 \ { m \AA} \ b &= & 6.782 \pm 1 \ { m \AA} \end{array}$
	$c = 5.747 \stackrel{\frown}{\pm} 1 \text{ Å}$
Cell volume	764.3 Å ³
Density (X-ray)	1.418 g cm^{-3}
Molecules per unit cell	2
Absent reflections	h00 with $h=2n+1$ and
	0k0 with $k=2n+1$
Space group	$P2_12_12$

The positional parameters obtained for the isomorphous strontium compound ¹ were used as starting parameters in a full matrix least-squares refinement of the calcium salt structure. The scattering factor curves used were derived from tables given by Freeman.⁷ The real parts of the corrections for anomalous dispersion were included. Attempted least-squares refinements with anisotropic temperature factors for atoms other than calcium did not converge.

Table 2. Final positional and thermal parameters with estimated standard deviations (σ).

			•	
Atom	$oldsymbol{x}$	$oldsymbol{y}$	\boldsymbol{z}	\boldsymbol{B}
Ca	0	0	0.2702 ± 8	
O(0)	0.0598 ± 6	0.171 ± 2	0.971 ± 3	2.2 ± 3
O(1)	0.1579 ± 7	$0.262~\overline{\pm}~2$	0.815 ± 3	$\textbf{2.8}\pm\textbf{3}$
O(2)	0.1059 ± 6	$\textbf{0.158} \pm \textbf{2}$	0.394 ± 2	1.5 ± 3
O(2')	0.2148 ± 7	0.947 ± 2	0.197 ± 3	$\textbf{2.9}\pm\textbf{3}$
O(4)	0.0889 ± 7	0.751 ± 2	0.160 ± 3	2.3 ± 3
O(5)	0.0285 ± 6	0.781 ± 2	0.585 ± 3	2.7 ± 3
C(1)	0.1203 ± 8	0.208 ± 3	0.985 ± 4	1.6 ± 4
C(2)	0.1529 ± 8	0.251 ± 3	0.225 ± 3	1.2 ± 3
C(2')	$\textbf{0.2227}\pm\textbf{8}$	$\boldsymbol{0.149\pm3}$	0.249 ± 4	1.4 ± 3
C(3)	0.1613 ± 9	0.471 ± 3	0.294 ± 3	1.9 ± 4
C(4)	0.0957 ± 10	0.595 ± 3	0.326 ± 4	$\textbf{2.3}\pm\textbf{4}$
C(5)	0.0886 ± 10	$\textbf{0.660}\pm\textbf{3}$	0.570 ± 4	$\textbf{2.3}\pm\textbf{4}$

For Ca the anisotropic temperature factor T obtained was $T = \exp[-0.0011 \ h^2 - 0.011 \ k^2 - 0.003 \ l^2 + 0.0001 \ hk]$

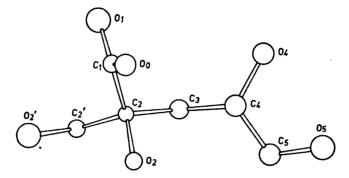


Fig. 1. The structure of the α-D-glucoisosaccharate group.

In the refinement Hughes' scheme ⁸ with $|F_{0,\,\mathrm{min}}|=4$ was used to assign weights to the 433 reflections used in the refinement. The measurements of the 26 weakest and 19 of the remaining intensities were obviously in error and these reflections were thus omitted from the least squares calculations. The final value of the residual $R[R=\sum||kF_{0}|-|F_{c}||/\sum|kF_{0}|]$ was 0.108. At this stage the shifts in all parameters were less than 1 % of their standard deviations. All attempts to refine the hydrogen positions were unsuccessful. The parameters obtained are listed in Table 2.

The structure of the α -D-glucoisosaccharate group (Fig. 1 and Table 3) is not significantly different from that found in the isomorphous strontium compound. Since the strontium compound is determined with higher precision

Table 3. Interatomic distances (in Å) and angles in the α -D-glucoisosaccharinic ion with standard deviations.

C(1) - C(2)	1.55 ± 3	O(0) - C(1) - C(2)	$119.9 + 1.6^{\circ}$
C(2) - C(2')	1.54 ± 2	O(1) - C(1) - C(2)	112.9 ± 1.4
C(2)-C(3)	1.55 ± 3	O(0) - C(1) - O(1)	125.0 ± 1.8
C(3)-C(4)	1.55 + 3	C(1) - C(2) - C(3)	116.9 + 1.4
C(4)-C(5)	1.47 + 3	C(1) - C(2) - C(2)	111.2 + 1.4
$C(\mathbf{x}) = C(0)$	1.11 ± 0		
		C(1) - C(2) - O(2)	104.3 ± 1.2
C(1) - O(0)	1.21 ± 2	C(2')-C(2)-C(3)	108.4 ± 1.3
C(1) - O(1)	1.28 ± 3	C(2')-C(2)-O(2)	107.7 + 1.3
C(2) - O(2)	1.48 + 2	O(2) - C(2) - C(3)	107.9 + 1.3
C(2') - O(2')	$1.41 \overset{-}{\pm} 2$	C(2)' - C(2') - O(2')	108.6 ± 1.3
C(4) - O(4)	1.43 ± 3	C(2) - C(3) - C(4)	117.7 ± 1.4
C(5) - O(5)	1.44 + 3	C(3) - C(4) - C(5)	110.8 ± 1.6
, , , ,	_	C(3) - C(4) - O(4)	$113.5 \stackrel{-}{\pm} 1.5$
O(0) - O(1)	2.21 + 2	O(4) - C(4) - C(5)	$113.9~\overset{-}{\pm}~1.6$
O(0) - O(2)	2.60 + 2	C(4) - C(5) - O(5)	107.8 + 1.6
O(2) - O(2')	$\frac{1}{2.81} + \frac{1}{2}$	- (-) - (-)	
O(4) - O(5)	2.72 ± 2		

(R = 0.058) we refer to the paper on the structure of Sr-3-deoxy-2-C-hydroxy-methyl-D-erythro-pentoate (i.e. Sr- α -D-glucoisosaccharate).¹

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The dimensions of the coordination polyhedron around each metal ion, a distorted Archimedean antiprism (Table 4), are somewhat decreased in the calcium compound. The average Me—O distance in the calcium salt is thus 0.12 Å less than in the strontium salt. As a result of this the O—O distances

Table 4. Ca-O distances (in Å) and intermolecular O-O distances (in Å) shorter than 3.2 Å. The O-O distances involved in the proposed hydrogen bonding scheme are marked with asterisks. The remaining O-O distances given in the table are all within the coordination polyhedron around calcium.

Ca - O(0) Ca - O(2) Ca - O(4) Ca - O(5)	$egin{array}{c} 2.38 \pm 2 \ 2.44 \pm 1 \ 2.51 \pm 1 \ 2.41 \pm 2 \ \end{array}$
O(0) - O(4) O(0) - O(4) * $O(0) - O(5)$ * $O(1) - O(2)$ * $O(1) - O(2')$ * $O(2') - O(4)$ O(2) - O(5) O(2) - O(5)	$egin{array}{c} 3.10 \pm 2 \ 3.15 \pm 2 \ 2.84 \pm 2 \ 2.72 \pm 2 \ 2.79 \pm 2 \ 2.81 \pm 2 \ 3.09 \pm 2 \ 2.88 \pm 2 \ 3.17 \pm 2 \ \end{array}$
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within the coordination polyhedron are somewhat less in the calcium salt. However, all short intermolecular O—O contacts outside the coordination polyhedron remain unchanged. This fact lends strong support to the proposed hydrogen bonding scheme.

EXTENDED HÜCKEL CALCULATIONS

The α -D-glucoisosaccharate ion shows two distinct structural features which will be briefly discussed here.

- 1. The intramolecular distance O(0)-O(2) is as short as 2.60 Å. A calculation of a least-squares plane through O(0), O(1), O(1), O(1), and O(2) shows that these atoms are without any significant deviation located in a plane. The atom O(2) is removed a very short distance, 0.30 Å, from this plane. This seems to be a common feature of all α -hydroxycarboxylic ions and acids. Interefore, the cation is probably not the cause of the short O(0)-O(2) distance.
- 2. The carbon chain is twisted at C(4) (cf. Fig. 1) and therefore only C(2'), C(2), C(3), and C(4) form a planar zigzag chain. This might be caused by the influence of the cation which is coordinated to O(4) and O(5), but it could also be associated with the fact that C(3) is the only carbon atom which is not bonded to an oxygen atom. The twist may be regarded as a rotation around C(3)—C(4). It may be pointed out that in the arabonate ion 12 and in the gluconate ion 12 all the carbon atoms are bonded to oxygen and form planar zigzag chains.

The shape of the anion is influenced by various forces in the crystal structure. Therefore it was considered to be of interest to make a comparison with the geometrical shape of a free acid molecule. By means of the extended Hückel program written by Hoffman ¹³ (rewritten by R. Piccioni and modified for CDC 3600 by G. Sperber) it was possible to make some semiempirical conformational analyses of the α -D-glucoisosaccharinic acid. A restriction which must be kept in mind is that the calculations apply to isolated, unsolvated acid molecules. The results obtained may therefore be of limited value but it must be remarked that there are very few ways to bridge the gap between a crystal structure and the corresponding molecular structure of a compound in solution.

It is beyond the scope of this paper to discuss the extended Hückel theory and we refer to the papers by Hoffmann.²⁻⁵

The following parameters were used for the calculations:

	Slater exponent	Coulombic integral (H_{ii})
$\mathbf{H}(1s)$	1.0	-13.60 eV
C(2s)	1.625	-21.34
C(2p)	1.625	11.54
O(2s)	2.275	-35.30
O(2p)	$\boldsymbol{2.275}$	-17.91

The resonance integrals H_{ij} (i \neq j) were approximated as $H_{ij}=0.5~K$ ($H_{ii}+H_{jj}$) S_{ij} with K=1.75 and where S_{ij} is the calculated overlap integral.² The structural parameters used were those of Sr- α -D-glucoisosaccharate. The atomic coordinates used in the calculations are given in Table 5. The

Table 5. Coordinates in Å for α -D-glucoisosaccharinic acid (from Sr- α -D-glucoisosaccharate) used in the extended Hückel calculations.

O(0)	-2.4177	0.1747	-1.3795
O(1)	-0.5315	-0.3467	-2.4636
O(2)	-1.3583	0.0002	0.9816
O(2')	-0.3130	-2.4440	0.0286
O(4)	0.0749	3.4660	-0.9945
O(5)	-0.7146	4.4675	1.4401
C(1)	-1.2013	-0.0794	-1.4129
C(2)	-0.4036	-0.0331	-0.0960
$\mathbf{C}(2')$	0.5064	-1.2694	0.0497
$\mathbf{C(3)}'$	0.5044	1.2087	-0.0628
C(4)	-0.2936	2.5368	0.0807
C(5)	-0.0044	3.1909	1.4235
$\mathbf{H}(0)$	-2.98	-0.75	-1.38
$\mathbf{H}(2)$	-0.73	-0.01	1.86
$\mathbf{H}(2')$	0.32	-3.31	0.03
$\mathbf{H}(4)$	-0.16	2.89	-1.87
$\mathbf{H}(5)$	-0.44	5.03	$\bf 2.32$
$\mathbf{H}(2'1)$	1.14	-1.27	0.93
$\mathbf{H}(2'2)$	1.14	-1.27	-0.83
$\mathbf{H}(3'1)$	1.31	-1.18	0.82
$\mathbf{H}(\mathbf{3'2})$	1.31	1.18	-0.94
$\mathbf{H}(4'1)$	-1.37	2.47	0.08
$\mathbf{H}(5'1)$	1.07	3.37	1.42
$\mathbf{H}(5'2)$	-0.26	2.62	2.30
11(0 2)	-0.20	2.02	2.30

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hydrogen atom positions were obtained from geometrical considerations assuming C—H distances of 1.1 Å and O—H distances of 1.0 Å. The calculated net charges and the overlap populations are shown in Figs. 2 and 3, respectively. They show why maxima were obtained in the electron density difference

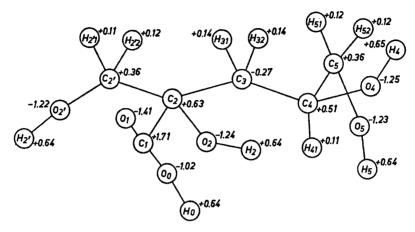


Fig. 2. Calculated net charges (electron units).

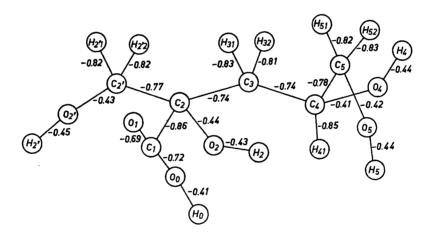


Fig. 3. Calculated overlap populations (electron units).

synthesis for hydrogen bonded to carbon but not for hydrogen bonded to oxygen. It is also evident from the net charges and overlap populations why X-ray analysis, when using atomic scattering functions, should give bond lengths which are too short for C—H and particularly short for O—H bonds.

It has been pointed out by Hoffmann ⁵ that the extended Hückel calculations are believed to be reliable if only the orientation of one part of a molecule

Table 6. Observed and calculated structure factors. Reflections marked with an asterisk were not used in the refinement. The columns give $h, k, l, |F_o|$ and $|F_c|$.

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503	469	46B				16 2 2	209	218	111	302 162 119	205
607	705	700	013	267 112	234 87				222	302	203
002	10	25	111	110	-27	023	295	225	,,,	162	156
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Table 6. Continued.

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is varied, or where the arguments are concerned with the electron distribution. Therefore, we have made some calculations for different conformations of the acid.

The carboxylic group was rotated around the C(1)-C(2) bond and the total energies calculated are plotted in Fig. 4. The minimum point is located about 18° from the position obtained in the crystal structure and corresponds to an almost planar group C(1), O(0), O(1), C(2), O(2). The deviation from a planar group is only 2°. Thus, the calculations show no tendency for the free acid to increase the short O(0)-O(2) distance by rotation of the carboxylic group. It should be stressed that the absolute values of the calculated energies are not to be relied upon. However, it seems probable that the energy involved in hydrogen bonding (in solution or in crystal structures) is too small to explain the configuration of the plane group.

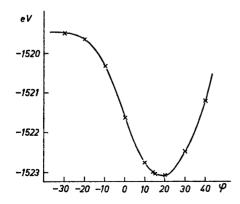


Fig. 4. Calculated total energy plotted as a function of rotation angle, φ , around the C(1)-C(2) bond. The φ -value 0° corresponds to the orientation found in the crystal structure.

Finally the C(3)—C(4) bond was rotated 75° in order to obtain a planar staggered zigzag chain. The calculated total energy was -1520.82 eV. This is 0.81 eV more than the energy calculated from the crystal structure parameters. Furthermore, it is notable that C(3) is the only carbon atom with a calculated negative net charge. Therefore, it is not improbable that the carbon chain remains twisted at C(4) in the solvated acid.

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