THE CRYSTAL STRUCTURE OF p-RIBONO-1,4-LACTONE AT -150°

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ABSTRACT

The crystal structure of p-ribono-1,4-lactone, 1, $C_5H_8O_5$, at -150° is orthorhombic, $P2_12_12_1$, with a=10.441(2), b=11.948(3), c=4.781(1) Å, V=596.43 Å (MoK α_1 , $\lambda=0.7107$ Å), Z=4, $D_m=1.626$ g cm⁻³ (20°), and $D_x=1.649$ g cm⁻³ (-150°). The 1,4-lactone ring has a conformation midway between E_3 and 2T_3 , with puckering comparable to that observed in the furanose rings of nucleosides. The C-O-C=O group has bond lengths of 1.467(1), 1.354(1), and 1.203(1) Å, respectively, and is slightly non-planar, with a C-O-C=O torsion angle of 172.7°. The molecules are linked by hydrogen bonds, which form chains. The carbonyl oxygen atom is involved in a weak, bifurcated hydrogen-bond interaction.

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INTRODUCTION

The crystal structures of six simple γ -lactones have been studied. These are this structure (1), D-gulono-1,4-lactone¹ (2), D-galactono-1,4-lactone² (3), 2,3,6-tri-O-methyl-D-galactono-1,4-lactone³ (4), D-glucaro-1,4-lactone⁴ (5), and β -D-glucurono-6,3-lactone⁵ (6).

Comparison of these results provides information relating to the geometrical characteristics of the γ -lactone ring system and to what degree this can be distorted by crystal-field effects.

EXPERIMENTAL

D-Ribono-1,4-lactone 1, D-ribonic γ -lactone, CAS No. 5336-08-3, from Vega Biochemicals, was recrystallized from 80% ethanol-water at room temperature. A crystal 0.31 \times 0.37 \times 0.55 mm was used to measure 2786 symmetry-independent intensities at -150° , using MoK α graphite-monochromated radiation on a CAD-4 diffractometer. Of these, 2418 had values $> 2\sigma_c$, based on counting statistics. No corrections were made for absorption ($\mu_{\text{MoK}\alpha} = 1.616$) or extinction. The unit-cell parameters were determined by least-squares fit of $\sin^2\theta$ values for 40 reflections with $36^\circ \le 2\theta \le 43^\circ$.

TABLE I FRACTIONAL ATOMIC COORDINATES a FOR D-RIBONO-1,4-LACTONE AT -150°

| | x | y | z | |
|-------|----------|---------------|-----------|--|
| C-1 | 33736(7) | 9747(6) | 32129(18) | |
| C-2 | 36756(7) | 15508(6) | 59463(17) | |
| C-3 | 51039(7) | 17848(6) | 56326(19) | |
| C-4 | 55262(7) | 7195(6) | 40785(19) | |
| C-5 | 58512(8) | -2578(7) | 59720(20) | |
| O-1 | 23781(6) | 9697(5) | 19457(15) | |
| O-2 | 28616(6) | 24733(5) | 63223(15) | |
| O-3 | 52498(6) | 27441(5) | 39194(19) | |
| O-4 | 44336(6) | 4396(5) | 22790(14) | |
| O-5 | 64710(6) | -11410(5) | 44691(16) | |
| H-2 | 356(1) | 102(1) | 734(4) | |
| H-3 | 558(1) | 186(1) | 723(4) | |
| H-4 | 623(1) | 86(1) | 280(3) | |
| H-51 | 507(1) | -55(1) | 688(3) | |
| H-52 | 648(1) | 2(1) | 735(3) | |
| H-O-2 | 306(2) | 288(1) | 773(4) | |
| H-O-3 | 598(2) | 287(1) | 364(4) | |
| H-O-5 | 590(2) | -144(1) | 356(4) | |

 $^{^{}a}$ Values are \times 10 5 for non-hydrogen atoms, \times 10 3 for hydrogen atoms. Estimated standard deviations given in parentheses refer to the least-significant digit.

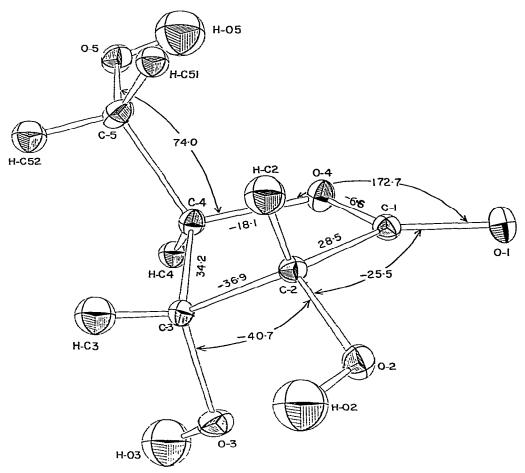


Fig. 1. D-Ribono-1,4-lactone at -150 °C. Atomic notation and thermal ellipsoids at 50% probability¹⁶. The numerical values refer to torsion angles (°). E.s.d. values are 0.1°.

The structure was solved with MULTAN⁶, using 106 reflections for phase generation. The first E-map showed all of the non-hydrogen atoms. The hydrogen atoms were located on difference maps at R=0.092. The parameter refinement was by full-matrix, least squares of $\sum \omega(|F_o|-|F_c|)^2$, where $\omega^{-1}=\sigma^2(F_o)+(0.02I)^2$, using ORFLS⁷ with anisotropic parameters for non-hydrogen atoms and isotropic thermal-parameters for hydrogen atoms. The final agreement factors were R=0.035, $R_\omega=0.037$, S=1.59, where $R=\sum ||F_o|-|F_c||/\sum |F_o|$, $R_\omega=[\sum \omega(|F_o|-|F_c|)^2/\sum |F_o|^2]^{1/2}$, and $S=(\sum \omega(|F_o|-|F_c|)/(m-n)^{1/2}$; m is the number of observations and n is the number of variable parameters. The atomic scattering-factors used for carbon and oxygen were those of Doyle and Turner⁸ and for hydrogen those of Stewart, Davidson, and Simpson⁹. The atomic parameters are given in Table I.

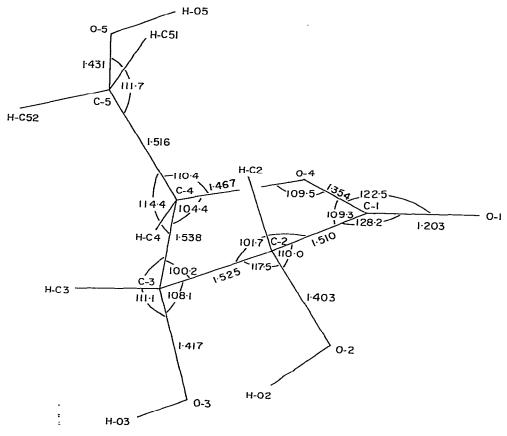


Fig. 2. Bond lengths (Å) and valence angles (°) in D-ribono-1,4-lactone at $-150\,^{\circ}$ C. E.s.d. values are 0.001 Å and 0.1°.

The atomic notation and thermal ellipsoids are shown in Fig. 1 and the molecular dimensions are given in Fig. 2*.

DISCUSSION

Because the (C-O-C=O) group is planar, or close to planar, the γ -lactone ring

is constrained to have a conformation 10 close to E_3 , this being less strained than the planar lactone ring. The data in Table II show ring conformations ranging from midway between 2T_3 ($\varphi = 90^\circ$) through E_3 ($\varphi = 108^\circ$) to midway to 4T_3 ($\varphi = 126^\circ$).

^{*}Tables of structure factors and anisotropic thermal-parameters are deposited with and can be obtained from: Elsevier Scientific Publishing Company, BBA Data Deposition, P. O. Box 1527, Amsterdam, The Netherlands. Reference should be made to No. BBA/DD/176/Carbohydr. Res., 92 (1981) 1-7.

The puckering ranges from q = 0.32 to 0.39 Å, a similar range to that observed in the furanose rings of nucleosides. The exception is 6, where the fused furanose ring constrains the lactone ring to an E_2 conformation ($\varphi = 252^{\circ}$) with less puckering. There are small deviations from planarity of the C-O-C=O group, as measured by the

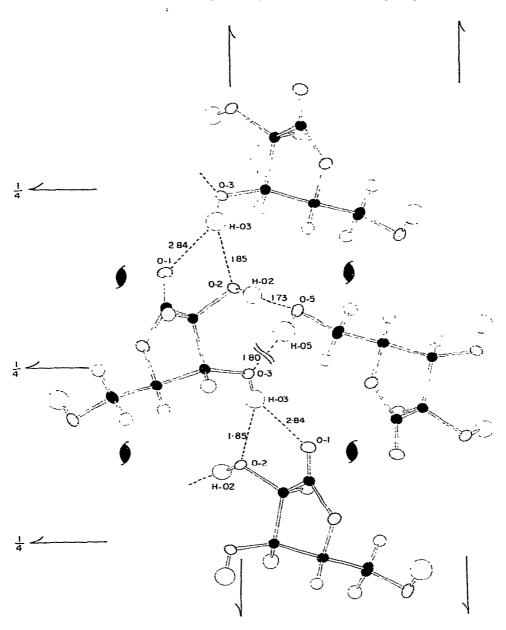


Fig. 3. Hydrogen bonding in the crystal structure of p-ribono-1,4-lactone. The values are $H\cdots O$ hydrogen-bond distances, where the O-H covalent-bond distances are normalized to 0.97 Å in the direction of the O-H bond.

| TABLE II |
|---|
| PUCKERING PARAMETERS ^a AND C–O BOND–LENGTHS FOR SOME 1,4-LACTONE RINGS |

| CRYST File name ^b | Puckering parameters | | Bond lengths (Å) | | | Torsion angle (deg) |
|---------------------------------|-------------------------|------|------------------|----------|----------|---------------------|
| | φ(deg) | q(A) | C-1-0-1 | C-1-0-4 | C-4-0-4 | C-4-0-4-C-1-0-1 |
| This structure | 99 | 0.38 | 1.203(1) | 1.354(1) | 1.467(1) | +172.7 |
| GULONO 2 | 107 | 0.38 | 1.194(3) | 1.346(3) | 1.473(3) | +178.4 |
| GALLAD 3 | 113 | 0.39 | 1.198(5) | 1.358(5) | 1.464(5) | -178.6 |
| MGALAC | 104 | 0.37 | 1.205(5) | 1.350(5) | 1.455(6) | +174.7 |
| DGLACM 5 | 102 | 0.32 | 1.210(5) | 1.337(4) | 1.466(4) | +175.4 |
| GLULAD 6 | 254 | 0.16 | 1.215(5) | 1.340(5) | 1.475(5) | -169.4 |

^aRef. 14. ^bThe six-letter acronym is the REFCODE for searching the Cambridge Crystallographic Data File¹⁵.

C-O-C=O torsion angles. These are presumably due to crystal-field forces, except in the case of 6, where a larger departure from planarity can be associated with the strain arising from the fused-ring structure. The C-O-C=O bond lengths show the characteristic lengthening and shortening of the formal C-O bonds because of contributions from the valence-bond resonance form $C^+-O=C-O^-$. The small differences in bond lengths are probably not significant, as they do not appear to be related in any systematic way to the small differences in conformation. The C-C bond lengths are normal, ranging from 1.510 to 1.538 Å, as are the C-OH bonds, 1.403 to 1.431 Å. The orientation of the primary alcohol group is gauche/trans.

The hydrogen-bonding scheme is shown as follows.

As in the crystal structure of 1,6-anhydro β -D-galactopyranose¹¹, the cooperative effect of the infinite chain of hydrogen bonds is maintained while the non-hydrogenated oxygen atom, O-1, is incorporated as a bifurcated interaction^{12,13}. The ring-oxygen atom, O-4, is not included in the hydrogen bonding; its nearest intermolecular neighbor is a methylene hydrogen atom (H-2) at 2.63 Å. The geometry of the hydrogen bonds is shown in Fig. 3. The infinite chain of hydrogen bonds forms a spiral that extends around the screw axis in the x direction. The separation of the two carbonyl oxygen atoms across the screw axes in a dipolar arrangement is such that the O-1···O-1 distance is 3.339(1) Å.

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