outre, l'angle $O \cdots C = O(\alpha)$ a une valeur proche de 100 à 110°. Enfin, l'atome O nucléophile montre une tendance à se placer dans le plan bissecteur de l'angle R_xCR_y . La plus courte distance intermoléculaire $O \cdots C$ qu'ils ont trouvée vaut 2,77 Å. Les valeurs des distances et des angles qui décrivent l'interaction $C \cdots O$ dans $C_6H_9NO_4$ sont en accord avec les conclusions de Bürgi *et al.*: si nous identifions R_x à O(71) et R_y à C(2), on trouve $\Delta = +0,011(2), \alpha =$ $96,6^\circ, \beta_3 (\equiv \angle R_xCO) = 90,8^\circ$ et $\beta_4 (\equiv \angle R_yCO) =$ $83,2^\circ$.

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The Crystal and Molecular Structure of 2-Acetamido-2,3-dideoxy-D-*erythro*-hex-2-enono-1,4-lactone, $C_8H_{11}NO_5$

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2-Acetamido-2,3-dideoxy-D-erythro-hex-2-enono-1,4-lactone crystallizes in space group $P2_12_12_1$ with a = 4.904 (1), b = 7.670 (1), c = 24.148 (3) Å, Z = 4. The structure was solved by direct methods and refined to an R of 0.058. The lactone ring is planar within experimental error. The molecules are linked in a three-dimensional network by intermolecular O(5)-H...O(7) (2.729), O(6)-H...O(5) (2.838) and N-H...O(1) (2.948 Å) hydrogen bonds.

Introduction

The crystal structure determination of 2-acetamido-2,3-dideoxy-D-*erythro*-hex-2-enono-1,4-lactone represents a continuation of our previous work on the structures of aminoacetylated unsaturated fivemembered-ring lactones. The crystal structure of the *threo* isomer has already been published (Ružić-Toroš & Kojić-Prodić, 1976). The inhibitory activities of 2-acetamido-2,3-dideoxy-D-hex-2-enonolactones on 2-acetamido-2-dideoxy- β -D-glucosidase were tested. Only the *erythro* isomers are inhibitors, whereas the *threo*

lactones are practically inactive. It was also observed that the 1,4-lactone inhibited 2-acetamido-2-deoxy- β -Dglucosidase more strongly than the 1,5-lactone (Pokorny, Zissis, Fletcher & Pravdić, 1975). The reason for the various inhibitory activities among these compounds was not obvious. It is very useful to make correlations between the molecular geometry and the inhibitory properties of these compounds and thus the crystal structure determination of the *erythro* isomer has been undertaken.

Experimental

Crystals of 2-acetamido-2,3-dideoxy-D-erythro-hex-2enono-1,4-lactone are colourless prisms. Unit-cell dimensions were determined from oscillation photographs with Cu $K\alpha$ radiation. The precise values (Table 1) were then deduced from zero-layer rotation patterns of single crystals (about the *a* and *c* axes) taken in asymmetric (Straumanis) positions and indexed by means of corresponding Weissenberg photographs (Popović, 1974).

Table 1. Crystallographic and physical data

2-Acetamido-2,3-dideoxy-D-*erythro*-hex-2-enono-1,4-lactone, $C_8H_{11}NO_5$

FW	201-18
Space group	P2,2,2,
0	4.904 (1) Å
u 1	7 (70 (1)
b	7·0/0(1)
С	24.148 (3)
U	908-3 Å ³
D,	1.472 g cm^{-3}
z	4
μ(Cu <i>K</i> α)	7.30 cm ⁻¹
Crystal shape	Prismatic
Crystal dimensions	$0.09 \times 0.18 \times 0.5$ mm
	424
F(000)	424

Table 2. Final positional parameters $(\times 10^4)$ for the non-hydrogen atoms

	x	у	Ζ
C(1)	-773 (12)	3200 (7)	606 (2)
C(2)	-333 (12)	1611 (7)	936 (2)
C(3)	-1851 (12)	1673 (7)	1390 (2)
C(4)	-3349 (12)	3368 (8)	1389 (2)
C(5)	-2651 (11)	4638 (7)	1852 (2)
C(6)	-3208 (13)	3945 (8)	2422 (3)
C(7)	1823 (12)	-1244 (7)	921 (2)
C(8)	3921 (15)	-2312 (8)	618 (3)
O(1)	265 (10)	3647 (5)	178 (2)
O(4)	-2650 (9)	4177 (5)	870 (2)
O(5)	189 (8)	5055 (5)	1824 (2)
O(6)	-5896 (10)	3281 (8)	2481 (2)
O(7)	521 (9)	-1769 (5)	1319 (2)
N	1484 (10)	389 (6)	720 (2)

The space group was determined from Weissenberg photographs recorded with Cu K radiation. Systematic absences indicated space group $P2_12_12_1$.

The intensities were collected on an Enraf-Nonius CAD-4 single-crystal automatic diffractometer with Nifiltered Cu $K\alpha$ radiation. Reflexions were scanned in the ω -2 θ mode (2 θ scan width = 0.6 + 0.2 tan θ). Intensities were measured in the range $1 < \theta < 90^{\circ}$. 2274 reflexions were collected from the *hkl* and *hkl* octants and were averaged to give a set of 1137 reflexions. 201 of these were unobserved ($I \le 2\sigma$). The data were corrected for variations in the intensity of the reference reflexions, and for Lorentz and polarization effects.

Structure determination and refinement

The structure was solved with MULTAN (Declerca, Germain, Main & Woolfson, 1973). Overall temperature and scale factors were determined (Wilson, 1942) and used to compute normalized structure amplitudes by the routine NORMAL included in MULTAN. The solution was based on 250 reflexions with $|E| \ge 1.2$. The E map corresponding to the solution with the best figure of merit (ABSFOM =1.31) revealed the positions of 11 non-hydrogen atoms. The positions of the other three non-hydrogen atoms were located from the resulting Fourier synthesis. A minimizing full-matrix least-squares procedure $\sum w ||F_o| - |F_c||^2$ with $w = 1/\hat{\sigma}_{|F_o|}^2$ was used for the refinement. Heavy-atom coordinates, isotropic thermal parameters and a scale factor were refined to an R of 0.118. H atoms were located from a difference Fourier synthesis after a few cycles of anisotropic refinement (R = 0.081). In the final cycle, one scale factor, the atomic positional parameters and the anisotropic thermal parameters for the non-hydrogen atoms were varied. For H atoms the coordinates were taken from a difference Fourier synthesis and the isotropic thermal parameters were those of the bonded atoms. The final

Table 3.	Positional	parameters	$(\times 10^{3})$	and	isotropic
therma	l parameter	$rs(\times 10^2)$ for	the hyd	rogen	atoms

	x	у	Z	U (Ų)
H(3)	-188	85	170	3.7
H(4)	-525	315	140	3.8
H(5)	-349	573	178	2.6
H(6,1)	-300	490	270	4.7
H(6,2)	-210	280	255	4.7
H(8,1)	320	-335	44	5.1
H(8.2)	420	-156	24	5.1
H(8.3)	520	-235	90	5.1
H(N)	190	60	38	3.2
HÌOS)	30	625	165	7.5
HÌOĜ	675	390	220	7.1

Table 4. Bond angles (°)

O(1)-C(1)-O(4)	122.5 (5)	O(4) - C(4) - H(4)	109
O(1)-C(1)-C(2)	129.8 (5)	C(3) - C(4) - H(4)	109
O(4)-C(1)-C(2)	107.7 (4)	C(5)-C(5)-H(4)	108
C(1)-C(2)-C(3)	109.6 (5)	O(5) - C(5) - C(4)	109.1 (5)
C(1)-C(2)-N	116.6 (5)	O(5) - C(5) - C(6)	107.5 (5)
C(3) - C(2) - N	133.8 (5)	C(4) - C(5) - C(6)	113.9 (5)
C(2)-C(3)-C(4)	107.8 (5)	O(6) - C(6) - C(5)	112.9 (5)
C(2)-C(3)-H(3)	128	O(7)–C(7)–C(8)	124.0 (5)
C(4) - C(3) - H(3)	124	O(7)–C(7)–N	121.5 (5)
O(4) - C(4) - C(3)	105.0 (5)	C(8)–C(7)–N	114.4 (5)
O(4) - C(4) - C(5)	108-1 (5)	C(2) - N - C(7)	124.5 (5)
C(3) - C(4) - C(5)	116.5(5)	C(1) - O(4) - C(4)	109.5(4)

agreement indices were $R = \Sigma ||F_o| - |F_c| / \Sigma |F_o| = 0.058$ and $R_w = [\Sigma w (|F_o| - |F_c|)^2 / |F_o|^2]^{1/2} =$ 0.069.

Scattering factors given by Cromer & Mann (1968) and (for H) by Stewart, Davidson & Simpson (1965) were used.

Table 5. Displacements from the least-squares plane (Å) for the lactone group

Atoms included in the calculation of the least-squares plane are denoted by an asterisk.

(2)*	-0.014
(1)*	-0.006
(1)*	-0.039
(4)*	0.025
(4)*	0.034
(3)	-0.013

C C O O C C

Table	6.	Torsion	angles	(°)	describing	the	confor-
	та	tions abo	ut C(4)-	-C($\overline{5}$) and $C(\overline{5})$ -	-C(6	5)

$\begin{array}{c} C(3)-C(4)-C(5)-O(5)\\ C(3)-C(4)-C(5)-H(5)\\ O(4)-C(4)-C(5)-O(5)\\ O(4)-C(4)-C(5)-C(6)\\ H(4)-C(4)-C(5)-C(6)\\ H(4)-C(4)-C(5)-H(5)\\ \end{array}$	59.4 (6) 171 -58.4 (6) -178.5 (5) 62 -66
$\begin{array}{c} C(4)-C(5)-C(6)-H(6,1)\\ C(4)-C(5)-C(6)-H(6,2)\\ O(5)-C(5)-C(6)-O(6)\\ O(5)-C(5)-C(6)-H(6,2)\\ H(5)-C(5)-C(6)-H(6,2)\\ H(5)-C(5)-C(6)-O(6)\\ H(5)-C(5)-C(6)-H(6,1)\\ \end{array}$	-171 62 -172·7 (5) -59 75 -44



Fig. 2. A view of the crystal structure along a showing the packing arrangement and hydrogen bonds.



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Table 7. Hydrogen bonds

$X - H \cdots Y$	$X \cdots Y$	X–H	$\mathbf{H}\cdots \mathbf{Y}$	$\angle X - H \cdots Y$	Symmetry operation
$O(5) - H(O5) \cdots O(7)$ $O(6) - H(O6) \cdots O(5)$	2·729 (6) Å 2·838 (7)	1.01 Å 0.93	1.72 Å 1.96	177° 156	x,y,z; x, y + 1, z
$N-H(N)\cdots O(1)$	2.948 (6)	0.86	2.21	144	$x,y,z; x + \frac{1}{2}, \frac{1}{2} - y, -z$

The calculations were carried out on the Univac 1110 computer at the University Computing Centre in Zagreb with the XRAY 72 system (Stewart, Kruger, Ammon, Dickinson & Hall, 1972). Positional parameters are listed in Tables 2 and 3.*

Description and discussion of the structure

The numbering of the atoms and the values of the bond lengths are shown in Fig. 1. The molecular packing and hydrogen bonds are illustrated in Fig. 2. Interatomic angles are listed in Table 4 and displacements of the atoms from the least-squares plane through the lactone group and ring are in Table 5. Dihedral angles defining the conformation about C(4)-C(5) and C(5)-C(6) are given in Table 6.

The atoms defining the lactone group are planar with a maximum deviation of 0.039 Å for O(1) (Table 5). The C(3) atom belonging to the lactone ring is displaced by 0.013 Å from the least-squares plane through the lactone group. Thus the lactone group and ring are both planar within the range of experimental error.

The conformation of the side chain about the C(4)-C(5) bond is 'staggered'; about the C(5)-C(6) bond the conformation is 'anti' (Table 6). The bond lengths and angles [with the exception of internal angles at C(2) and C(3)] are comparable with the values for the relevant type of hybridization, and with those found in L-ascorbic acid (Hvoslef, 1968), and the *threo* isomer of this compound (Ružić-Toroš & Kojić-Prodić, 1976).

The C-C bond lengths are in the range 1.472-1.522Å. The value of C(2)=C(3) [1.325 (8) Å] is equal to that in the *threo* isomer of this compound (Ružić-Toroš & Kojić-Prodić, 1976) and is comparable with 1.337 (6) Å (*International Tables for X-ray Crystallography*, 1968). The single bonds C(1)-C(2) [1.472 (8)] and C(3)-C(4) [1.494 (8) Å] are shortened compared with the standard value of 1.54 Å, owing to the influence of the C=O and C=C double bonds. However, they are normal for this structure. The asymmetry of the endocyclic C(1)–O(4) [1.347(7)]and C(4)–O(4) [1.440(7) Å] bonds observed in the present structure was also found in L-ascorbic acid (Hvoslef, 1968), γ -D-gulonolactone (Berman, Rosenstein & Southwick, 1971) and the *threo* isomer of this compound (Ružić-Toroš & Kojić-Prodić, 1976). This asymmetry can be interpreted by the valence resonance form $-C=C^+-C-$. Other single C–O bonds of \parallel O⁻

1.421 (8) and 1.431 (7) Å, and double C=O bonds of 1.201 (7) and 1.222 (7) Å are in the usual range. The C-N bonds are 1.353 (7) and 1.395 (7) Å.

The internal angles C-C-C, C-O-C and O-C-C are in the range $105 \cdot 0$ (4)- $109 \cdot 6$ (5)°. The mean value of $107 \cdot 9^{\circ}$ is nearly equal to the calculated angle of 108° in a regular pentagon. The influence of fivemembered-ring geometry on the values of bond angles in the lactone ring at C(1) $[107 \cdot 7 (4)^{\circ}]$, C(2) $[109 \cdot 6 (5)]$ and C(3) $[107 \cdot 8 (5)^{\circ}]$ is obvious. The same distortions are noticed in L-ascorbic acid (Hvoslef, 1968) and the *threo* isomer (Ružić-Toroš & Kojić-Prodić, 1976).

The molecules are connected by hydrogen bonds forming a three-dimensional network (Fig. 2 and Table 7). Both hydroxyl groups are involved in hydrogen bonds. The O(5) hydroxyl group acts as an acceptor in O(6)-H(O6)...O(5) [2.838 (7) Å] and as a donor to a carbonyl O atom in the O(5)-H(O5)...,O(7) [2.729 (6) Å] hydrogen bond. The N atom in the acetamido group acts as a donor to the second carbonyl O forming the N-H...O(1) [2.948 (6) Å] hydrogen bond.

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^{*} Lists of structure factors and anisotropic thermal parameters have been deposited with the British Library Lending Division as Supplementary Publication No. SUP 33064 (11 pp.). Copies may be obtained through The Executive Secretary, International Union of Crystallography, 13 White Friars, Chester CH1 1NZ, England.

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Stereochemistry of Unsaturated Amino Sugars. III. The Crystal and Molecular Structure of Peracetylated 1,2-Dideoxy-D-arabinoaldopyranose, C₁₆H₂₁O₉N

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3,4,6-Tri-O-acetyl-2-(N-acetylacetamido)-1,2-dideoxy-D-arabino-hex-1-enopyranose crystallizes in space group $P2_12_12$ with a = 11.805, b = 18.586, c = 8.3196 Å, Z = 4. The structure was solved by MULTAN and refined with a full-matrix least-squares procedure to R = 0.045. The sugar ring exhibits an alternate sofa conformation; four C atoms and the ring O atom are coplanar and only one C atom is below this plane. The substituents are attached to C(3) in quasi-equatorial, and to C(4) and C(5) in equatorial positions. Owing to the absence of free hydroxyl groups, hydrogen bonds of the type $O-H\cdots O$ are not possible. In the crystal lattice the molecules are joined by van der Waals interactions only.

Introduction

3,4,6-Tri-O-acetyl-2-(N-acetylacetamido)-1,2-dideoxy-D-arabino-hex-1-enopyranose was synthesized (Pravdić & Fletcher, 1967; Pravdić, Franjić-Mihalić & Danilov, 1975) as the first representative of a new class of unsaturated amino sugars. The crystal structures of three compounds from this series have already been solved (Kojić-Prodić, Rogić & Ružić-Toroš, 1976; Rogić, Ružić-Toroš, Kojić-Prodić & Pravdić, 1977). The main interest of the structural study of the above series is the sugar-ring conformation. The presence of the double bond in these ring systems suggests a halfchair conformation. Indeed, it was found to be highly predominant in this series. The present structure determination, as part of this study, shows the sugar ring to be in an unusual alternate sofa conformation.

Experimental

The space group was determined as $P2_12_12$ from Weissenberg photographs recorded with Cu Ka

radiation. Cell constants were obtained by the leastsquares procedure using the θ values for 19 reflexions in the interval 50.2° < 2θ < 57.6° (at 18°C) with Cu K α radiation. Crystal data are given in Table 1. Integrated intensities for 2151 independent reflexions with θ < 75° were measured on an Enraf–Nonius CAD-4 diffractometer using Cu K α radiation. The data were corrected for background, Lorentz and polarization effects.

Structure determination and refinement

The structure was solved by *MULTAN* (Germain, Main & Woolfson, 1971) using normalized structure amplitudes calculated by the *K*-curve method (Hauptman & Karle, 1953). The solution was based on 271 reflexions with $|E| \ge 1.50$. The *E* map corresponding to the solution with the best figure of merit revealed the positions of all the non-hydrogen atoms. The H atoms were located from a difference Fourier map with the exception of H(8,2), H(10,3), H(12,3), and H(16,3). Their positions were calculated on stereo-

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