Acta Crystallographica Section E

Structure Reports Online

ISSN 1600-5368

2C-Methyl-D-arabinono-1,4-lactone monohydrate

Francesco Punzo, ** David J. Watkin, * Sarah F. Jenkinson and George W. J. Fleet

^aDipartimento di Scienze Chimiche, Facoltà di Farmacia, Università di Catania, Viale A. Doria 6, 95125 Catania, Italy, ^bDepartment of Chemical Crystallography, Chemical Research Laboratory, Mansfield Road, Oxford OX1 3TA, England, and ^cDepartment of Organic Chemistry, Chemical Research Laboratory, Mansfield Road, Oxford OX1 3TA, England

‡ Visiting Scientist at the Department of Chemical Crystallography, Chemical Research Laboratory, Mansfield Road, Oxford OX1 3TA, England.

Correspondence e-mail: francesco.punzo@chemistry.oxford.ac.uk

Key indicators

Single-crystal X-ray study T = 120 KMean $\sigma(\text{C-C}) = 0.002 \text{ Å}$ R factor = 0.031 wR factor = 0.071Data-to-parameter ratio = 12.5

For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e.

The title compound, $C_6H_{10}O_5 \cdot H_2O$, formed by the hydrolysis of a δ -lactone, is shown unequivocally to be a γ -lactone. The diol has a *trans* configuration.

Received 10 December 2004 Accepted 10 January 2005 Online 22 January 2005

Comment

The potential of the Kiliani ascension of ketoses to provide readily available branched scaffolds has been recognized (Hotchkiss et al., 2004). A further class of branched carbohydrate building blocks may be available from the reaction of cyanide on 1-deoxyketoses, themselves prepared by addition of organometallic reagents to sugar lactones. The protected 1deoxy-D-ribulose, (1), was treated with sodium cyanide and gave a single diastereomeric product, (2), the structure of which was established by X-ray crystallography (Punzo et al., 2005). During the isolation of (2), some loss of the protecting acetonide group afforded an unprotected lactone (3), which was eventually crystallized. NMR and other structural studies on (3) could not firmly determine the size of the lactone ring; X-ray crystallographic analysis established that (3) is a 1,4lactone (Fig. 1). It is noteworthy that none of the epimeric ribonolactone, (4), was isolated during the course of the synthesis. As usually expected for sugar derivatives, hydrogen bonding (Table 2) occurs between molecules, and the water of crystallization is involved in this network (Fig. 2).

Experimental

Compound (3) was crystallized by dissolving it in diethyl ether, adding a few drops of cyclohexane and allowing the slow competitive evaporation of the two solvents until clear colourless crystals formed. Water was used as solvent during the synthesis of the compound. Moreover the compound was exposed to air before and after crystallization.

doi:10.1107/S1600536805000723

Crystal data

 $C_6H_{10}O_5 \cdot H_2O$ $M_r = 180.16$ Orthorhombic, $P2_12_12_1$ a = 8.1624 (3) Å b = 8.5569 (3) Å c = 11.6000 (5) Å V = 810.20 (5) Å³ Z = 4 $D_x = 1.477 \text{ Mg m}^{-3}$ Mo K α radiation Cell parameters from 1300 reflections $\theta = 5-30^{\circ}$ $\mu = 0.13 \text{ mm}^{-1}$ T = 120 KPlate, colourless $0.30 \times 0.20 \times 0.04 \text{ mm}$

© 2005 International Union of Crystallography Printed in Great Britain – all rights reserved

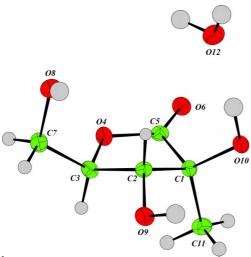


Figure 1
The asymmetric unit of (3), with displacement ellipsoids drawn at the 50% probability level.

Data collection

Nonius KappaCCD diffractometer ω scans Absorption correction: multi-scan (DENZO/SCALEPACK; Otwinowski & Minor, 1997) $T_{\min} = 0.97, T_{\max} = 0.99$ 2296 measured reflections 1361 independent reflections 1201 reflections with $I > 2 \sigma(I)$ $R_{\rm int} = 0.013$ $\theta_{\rm max} = 30.0^{\circ}$ $h = -11 \rightarrow 11$ $k = -11 \rightarrow 12$ $l = -16 \rightarrow 16$

Refinement

Refinement on F^2 $R[F^2 > 2\sigma(F^2)] = 0.031$ $wR(F^2) = 0.071$ S = 0.981361 reflections 109 parameters H-atom parameters constrained
$$\begin{split} w &= 1/[\sigma^2(F^2) + 0.03 + 0.17P], \\ \text{where } P &= [\max(F_o^2, 0) + 2F_c^2]/3 \\ (\Delta/\sigma)_{\max} &< 0.001 \\ \Delta\rho_{\max} &= 0.23 \text{ e Å}^{-3} \\ \Delta\rho_{\min} &= -0.24 \text{ e Å}^{-3} \\ \text{Absolute structure: see text} \end{split}$$

Table 1
Selected bond lengths (Å).

C1-C2	1.537 (2)	C3-O4	1.4695 (18)
C1-C5	1.528 (2)	C3-C7	1.516(2)
C1-O10	1.4169 (17)	O4-C5	1.3363 (18)
C1-C11	1.526 (2)	C5-O6	1.2106 (17)
C2-C3	1.525 (2)	C7-O8	1.4261 (19)
$C_{2}^{2} = O_{9}$	1 4167 (18)		` ′

Table 2 Hydrogen-bonding geometry (Å, °).

D-H··· A	<i>D</i> —Н	$H \cdot \cdot \cdot A$	$D \cdot \cdot \cdot A$	$D-H\cdots A$
O10-H5···O12 O8-H7···O6 ⁱ O9-H9···O8 ⁱ O12-H12···O9 ⁱⁱ	0.92 0.97 0.96 0.94	1.81 1.78 1.76 2.01	2.7191 (16) 2.7235 (15) 2.7157 (15) 2.9138 (16)	175 163 169 163
$O12-H1\cdots O10^{iii}$	0.91	2.00	2.8613 (16)	157

Symmetry codes: (i) $1 - x, \frac{1}{2} + y, \frac{3}{2} - z$; (ii) $1 - x, y - \frac{1}{2}, \frac{3}{2} - z$; (iii) $x - \frac{1}{2}, \frac{1}{2} - y, 1 - z$.

In the absence of significant anomalous scattering, Friedel pairs were merged. The absolute configuration was assigned since the starting material was D-erythronolactone with known absolute

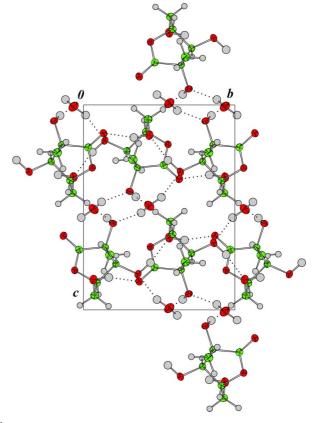


Figure 2 Packing diagram of (3), viewed down the *a* axis. Hydrogen bonds are indicated by dashed lines.

configuration. H atoms were located in difference density maps. Those attached to C atoms were repositioned geometrically. The H atoms were initially refined with soft restraints on the bond lengths and angles to regularize their geometry (C—H = 0.97–1.01 Å and O—H = 0.91–0.97 Å), after which they were refined as riding, with $U_{\rm iso}({\rm H})=1.2U_{\rm eq}({\rm C})$ and $U_{\rm iso}({\rm H})=0.05$ Ų for those bonded to the O atoms

Data collection: *COLLECT* (Nonius, 2001); cell refinement: *DENZO/SCALEPACK* (Otwinowski & Minor, 1997); data reduction: *DENZO/SCALEPACK*; program(s) used to solve structure: *SIR*92 (Altomare *et al.*, 1994); program(s) used to refine structure: *CRYSTALS* (Betteridge *et al.*, 2003); molecular graphics: *CAMERON* (Watkin *et al.*, 1996); software used to prepare material for publication: *CRYSTALS*.

References

Altomare, A., Cascarano, G., Giacovazzo, C., Guagliardi, A., Burla, M. C., Polidori, G. & Camalli, M. (1994). *J. Appl. Cryst.* 27, 435.

Betteridge, P. W., Carruthers, J. R., Cooper, R. I., Prout, K. & Watkin, D. J. (2003). J. Appl. Cryst. 36, 1487.

Hotchkiss, D., Soengas, R., Simone, M. I., van Ameijde, J., Hunter, S., Cowley, A. R. & Fleet, G. W. J. (2004). *Tetrahedron Lett.* **45**, 9461–9464.

Nonius (2001). COLLECT. Nonius BV, Delft, The Netherlands.

Otwinowski, Z. & Minor, W. (1997). *Methods in Enzymology*, Vol. 276, *Macromolecular Crystallography*, Part A, edited by C. W. Carter Jr & R. M. Sweet, pp. 307–326. New York: Academic Press.

Punzo, F., Watkin, D. J., Jenkinson, S. F. & Fleet, G. W. J. (2005). Acta Cryst. E61, 0127–0129.

Watkin, D. J., Prout, C. K. & Pearce, L. J. (1996). CAMERON. Chemical Crystallography Laboratory, Oxford, England.