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Key indicators

Single-crystal X-ray study
 $T = 190\text{ K}$
 Mean $\sigma(\text{C-C}) = 0.003\text{ \AA}$
 R factor = 0.037
 wR factor = 0.092
 Data-to-parameter ratio =

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

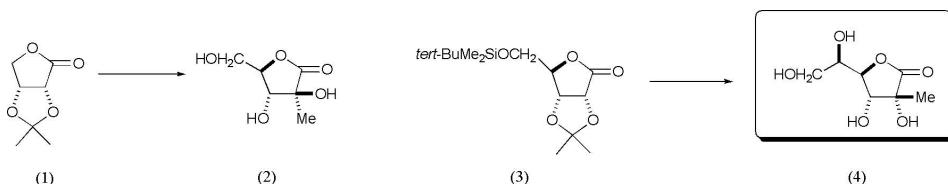
2-C-Methyl-D-allono-1,4-lactone

The relative stereochemistry at the quaternary C atom in the title compound, $C_7H_{12}O_6$, a 1,4-lactone formed from a protected D-ribonolactone, is firmly established by X-ray crystallographic analysis.

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Comment

The potential of the Kiliani ascension of ketoses to provide readily available branched scaffolds has been recognized (Harding *et al.*, 2005; Hotchkiss *et al.*, 2004; Shallard-Brown *et al.*, 2004); such materials are likely to be of value as a new family of chirons. A further class of branched carbohydrate building blocks may be available from the reaction of cyanide with 1-deoxyketoses, themselves prepared by addition of organometallic reagents to sugar lactones.



For example, when such a sequence was performed on the acetonide of D-erythronolactone (1), 2-C-methyl-D-arabinonolactone (2) was formed (Punzo *et al.*, 2005), in which the 2,3-diol unit is *trans*; none of the epimeric ribonolactone was isolated during the course of the synthesis. When a similar synthetic sequence was applied to the protected D-ribonolactone (3), the crystalline product (4) was isolated, in which the 2,3-diol unit is *cis*. There is no reliable spectroscopic technique to establish the relative stereochemistry in (4), so its relative configuration was unambiguously defined by X-ray crystallographic analysis; the absolute stereochemistry was defined by the use of D-ribonolactone as a starting material.

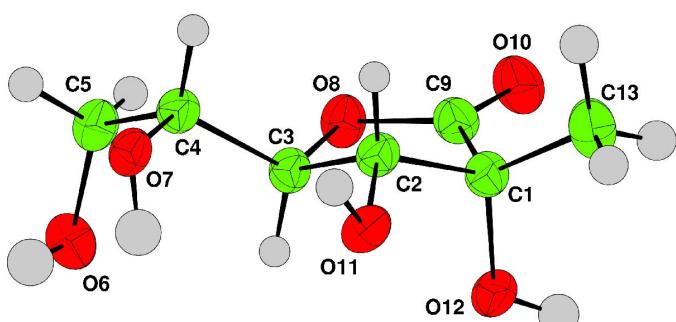
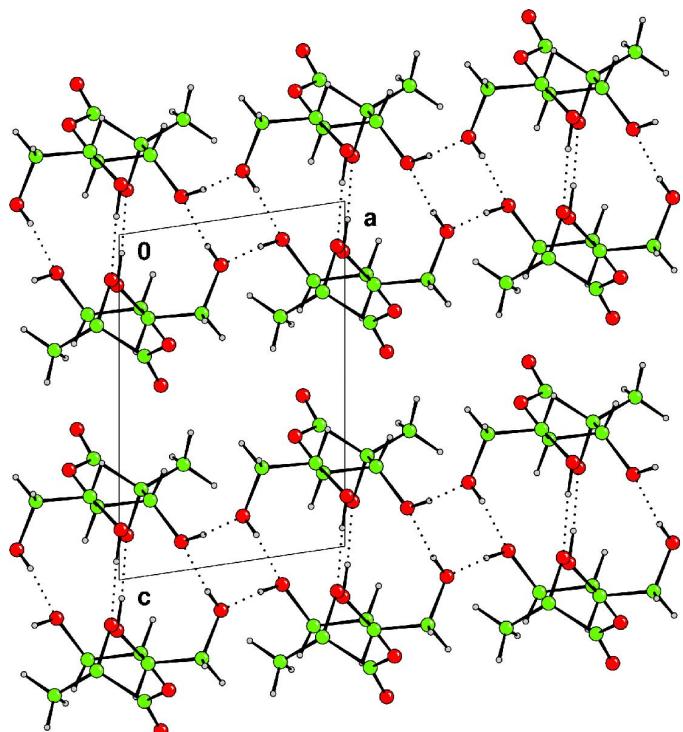
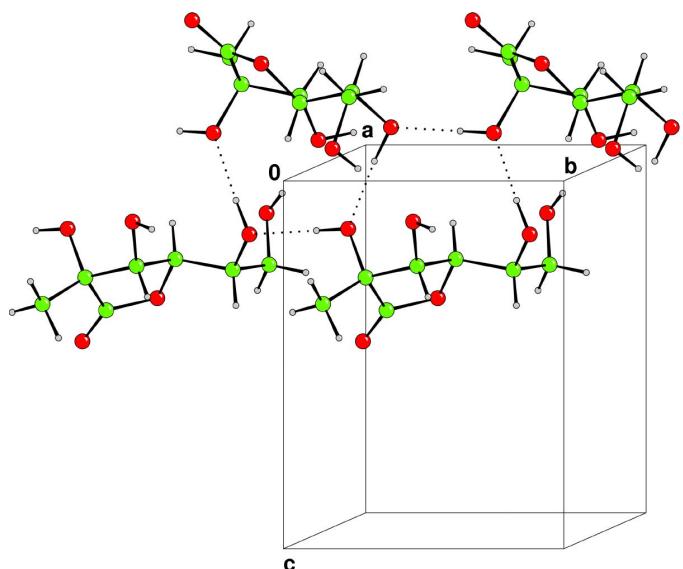


Figure 1

The molecular structure of (4), with displacement ellipsoids drawn at the 50% probability level.

**Figure 2**

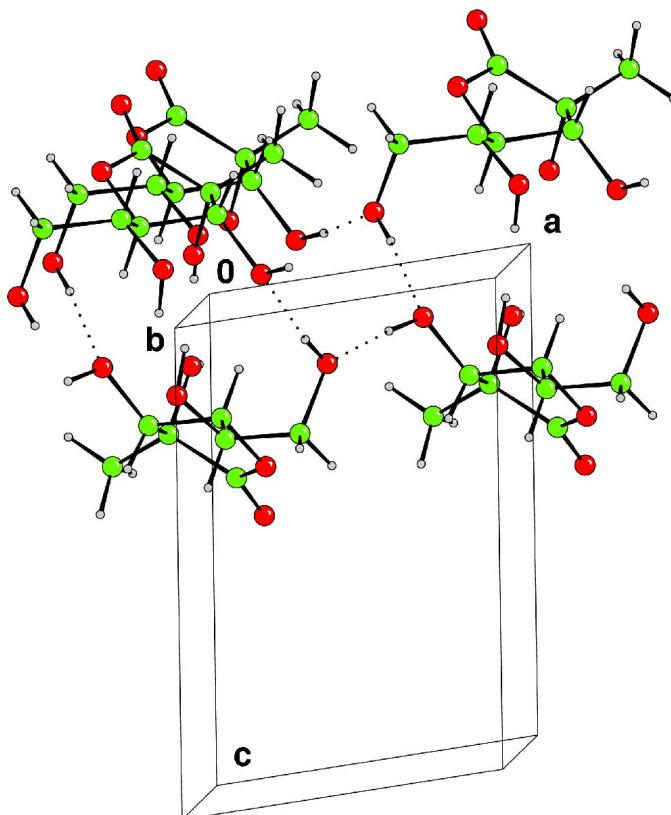
Packing diagram for the title compound, viewed down the *b* axis. Dashed lines indicate hydrogen bonds.

**Figure 3**

Hydrogen bonding (dashed lines): the zigzag network, forming columns of molecules.

The reactions are being studied further in order to understand the difference in the stereochemical outcome of the two sequences.

The crystal structure is made up of layers of strongly hydrogen-bonded molecules which lie in the *ab* plane. The layers are made up of columns of molecules along the *b* axis held together by a zigzag chain of hydrogen bonds, which are in turn tied together by a helical hydrogen-bonding network (Figs. 3 and 4).

**Figure 4**

Hydrogen bonding (dashed lines): the helical network which links the columns of molecules together to form a sheet.

Experimental

Crystals of the title compound were obtained by evaporation of a solution in an ethyl acetate/cyclohexane mixture, yielding colourless crystals. The full synthetic procedure will be published separately (Jenkinson *et al.*, 2005).

Crystal data

$C_7H_{12}O_6$	$D_x = 1.493 \text{ Mg m}^{-3}$
$M_r = 192.17$	Mo $K\alpha$ radiation
Monoclinic, $P2_1$	Cell parameters from 1283 reflections
$a = 6.1521 (5) \text{ \AA}$	$\theta = 5\text{--}27^\circ$
$b = 7.5495 (7) \text{ \AA}$	$\mu = 0.13 \text{ mm}^{-1}$
$c = 9.3055 (8) \text{ \AA}$	$T = 190 \text{ K}$
$\beta = 98.501 (5)^\circ$	Block, colourless
$V = 427.45 (6) \text{ \AA}^3$	$0.20 \times 0.10 \times 0.10 \text{ mm}$
$Z = 2$	

Data collection

Nonius KappaCCD diffractometer	$R_{\text{int}} = 0.026$
ω scans	$\theta_{\text{max}} = 27.5^\circ$
Absorption correction: none	$h = -7 \rightarrow 7$
2940 measured reflections	$k = -9 \rightarrow 8$
1025 independent reflections	$l = -11 \rightarrow 12$
905 reflections with $I > 2\sigma(I)$	

Refinement

Refinement on F^2	H-atom parameters constrained
$R[F^2 > 2\sigma(F^2)] = 0.037$	$w = 1/[\sigma^2(F^2) + (0.04P)^2]$,
$wR(F^2) = 0.092$	where $P = [\max(F_o^2, 0) + 2F_c^2]/3$
$S = 1.05$	$(\Delta/\sigma)_{\text{max}} < 0.001$
1025 reflections	$\Delta\rho_{\text{max}} = 0.27 \text{ e \AA}^{-3}$
118 parameters	$\Delta\rho_{\text{min}} = -0.22 \text{ e \AA}^{-3}$

Table 1
Hydrogen-bonding geometry (\AA , $^\circ$).

$D-\text{H}\cdots A$	$D-\text{H}$	$\text{H}\cdots A$	$D\cdots A$	$D-\text{H}\cdots A$
O12—H4 \cdots O7 ⁱ	0.88	1.84	2.723 (2)	178
O11—H6 \cdots O6 ⁱⁱ	0.96	1.73	2.681 (2)	175
O6—H8 \cdots O11 ⁱⁱⁱ	0.83	1.95	2.778 (2)	174
O7—H14 \cdots O12 ⁱⁱⁱ	0.96	1.84	2.791 (2)	175

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

All H atoms were observed in a difference electron density map and were refined using slack restraints to optimize their geometry [$\text{C}-\text{H} = 0.98 \text{ \AA}$, $\text{O}-\text{H} = 0.82 \text{ \AA}$, with $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$ or $1.5U_{\text{eq}}(\text{O})$], then made to ride on their parent atoms. In the absence of significant anomalous scattering effects, Friedel pairs were merged. The absolute configuration is known from the synthesis.

Data collection: *COLLECT* (Nonius, 1997); cell refinement: *DENZO/SCALEPACK* (Otwinowski & Minor, 1997); data reduction: *DENZO/SCALEPACK*; program(s) used to solve structure: *SIR92* (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*.

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supporting information

Acta Cryst. (2005). E61, o1472–o1474 [https://doi.org/10.1107/S1600536805012328]

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 $V = 427.45 (6)$ Å³
 $Z = 2$

$F(000) = 204$
 $D_x = 1.493 \text{ Mg m}^{-3}$
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 Cell parameters from 1283 reflections
 $\theta = 5\text{--}27^\circ$
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 Block, colourless
 $0.20 \times 0.10 \times 0.10$ mm

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Nonius KappaCCD
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 2940 measured reflections
 1025 independent reflections

905 reflections with $I > 2\sigma(I)$
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 $\theta_{\text{max}} = 27.5^\circ, \theta_{\text{min}} = 5.2^\circ$
 $h = -7 \rightarrow 7$
 $k = -9 \rightarrow 8$
 $l = -11 \rightarrow 12$

Refinement

Refinement on F^2
 Least-squares matrix: full
 $R[F^2 > 2\sigma(F^2)] = 0.037$
 $wR(F^2) = 0.092$
 $S = 1.05$
 1025 reflections
 118 parameters
 1 restraint

Primary atom site location: structure-invariant
 direct methods
 Hydrogen site location: inferred from
 neighbouring sites
 H-atom parameters constrained
 $w = 1/[\sigma^2(F^2) + (0.04P)^2]$,
 where $P = [\max(F_o^2, 0) + 2F_c^2]/3$
 $(\Delta/\sigma)_{\text{max}} = 0.000076$
 $\Delta\rho_{\text{max}} = 0.27 \text{ e } \text{\AA}^{-3}$
 $\Delta\rho_{\text{min}} = -0.22 \text{ e } \text{\AA}^{-3}$

Fractional atomic coordinates and isotropic or equivalent isotropic displacement parameters (Å²)

	x	y	z	$U_{\text{iso}}^*/U_{\text{eq}}$
C1	-0.0984 (3)	0.3222 (3)	0.2531 (2)	0.0320
C2	-0.1385 (4)	0.5174 (3)	0.2177 (2)	0.0296
C3	0.0955 (3)	0.5864 (3)	0.2218 (2)	0.0290
C4	0.1288 (3)	0.7823 (3)	0.2541 (2)	0.0304

C5	0.3673 (3)	0.8415 (3)	0.2636 (3)	0.0357
O6	0.4517 (2)	0.8076 (2)	0.13170 (17)	0.0386
O7	-0.0091 (3)	0.8810 (2)	0.14479 (17)	0.0312
O8	0.2203 (2)	0.4860 (2)	0.34013 (17)	0.0345
C9	0.1124 (3)	0.3352 (3)	0.3624 (2)	0.0324
O10	0.1853 (3)	0.2291 (3)	0.45428 (18)	0.0440
O11	-0.2766 (2)	0.5436 (2)	0.08380 (15)	0.0339
O12	-0.0337 (3)	0.2406 (2)	0.12701 (17)	0.0373
C13	-0.2852 (4)	0.2248 (4)	0.3069 (3)	0.0449
H21	-0.1993	0.5739	0.2957	0.0336*
H31	0.1497	0.5606	0.1295	0.0337*
H41	0.0760	0.8081	0.3461	0.0353*
H51	0.3799	0.9682	0.2861	0.0430*
H52	0.4571	0.7735	0.3426	0.0409*
H131	-0.2364	0.1024	0.3341	0.0549*
H132	-0.3180	0.2916	0.3959	0.0531*
H133	-0.4211	0.2216	0.2324	0.0543*
H4	-0.0218	0.1243	0.1329	0.0585*
H6	-0.3733	0.6401	0.0949	0.0487*
H8	0.3924	0.8804	0.0718	0.0768*
H14	0.0116	0.8276	0.0546	0.0746*

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
C1	0.0369 (11)	0.0294 (12)	0.0292 (10)	-0.0010 (10)	0.0033 (8)	0.0017 (10)
C2	0.0316 (10)	0.0281 (13)	0.0281 (10)	-0.0016 (9)	0.0013 (8)	-0.0015 (9)
C3	0.0320 (11)	0.0250 (12)	0.0296 (10)	0.0002 (9)	0.0030 (8)	0.0013 (9)
C4	0.0349 (11)	0.0269 (12)	0.0290 (10)	-0.0001 (9)	0.0031 (8)	0.0004 (8)
C5	0.0342 (11)	0.0322 (12)	0.0394 (11)	-0.0033 (10)	0.0015 (9)	-0.0013 (10)
O6	0.0329 (8)	0.0388 (10)	0.0449 (9)	0.0061 (8)	0.0085 (7)	0.0091 (8)
O7	0.0344 (7)	0.0237 (8)	0.0345 (8)	0.0016 (6)	0.0020 (6)	0.0003 (7)
O8	0.0342 (8)	0.0295 (9)	0.0381 (8)	0.0004 (7)	-0.0006 (7)	0.0044 (7)
C9	0.0351 (11)	0.0317 (12)	0.0305 (11)	0.0008 (10)	0.0055 (8)	0.0029 (9)
O10	0.0488 (9)	0.0425 (11)	0.0396 (9)	0.0022 (9)	0.0025 (7)	0.0134 (8)
O11	0.0378 (8)	0.0290 (9)	0.0324 (7)	0.0044 (7)	-0.0036 (6)	-0.0028 (7)
O12	0.0527 (10)	0.0240 (8)	0.0354 (9)	0.0027 (8)	0.0070 (7)	-0.0009 (7)
C13	0.0451 (13)	0.0413 (15)	0.0475 (14)	-0.0076 (12)	0.0043 (11)	0.0082 (12)

Geometric parameters (\AA , ^\circ)

C1—C2	1.522 (3)	C5—O6	1.425 (3)
C1—C9	1.528 (3)	C5—H51	0.980
C1—O12	1.432 (3)	C5—H52	0.994
C1—C13	1.511 (4)	O6—H8	0.828
C2—C3	1.526 (3)	O7—H14	0.956
C2—O11	1.415 (2)	O8—C9	1.349 (3)
C2—H21	0.964	C9—O10	1.208 (3)

C3—C4	1.517 (3)	O11—H6	0.956
C3—O8	1.459 (2)	O12—H4	0.882
C3—H31	0.986	C13—H131	0.993
C4—C5	1.524 (3)	C13—H132	1.015
C4—O7	1.433 (3)	C13—H133	1.004
C4—H41	0.979		
C2—C1—C9	100.2 (2)	C5—C4—H41	109.4
C2—C1—O12	107.2 (2)	O7—C4—H41	106.2
C9—C1—O12	105.2 (2)	C4—C5—O6	111.6 (2)
C2—C1—C13	115.8 (2)	C4—C5—H51	110.1
C9—C1—C13	115.0 (2)	O6—C5—H51	109.4
O12—C1—C13	112.3 (2)	C4—C5—H52	108.4
C1—C2—C3	101.7 (2)	O6—C5—H52	108.1
C1—C2—O11	112.5 (2)	H51—C5—H52	109.2
C3—C2—O11	114.6 (2)	C5—O6—H8	106.3
C1—C2—H21	109.5	C4—O7—H14	105.2
C3—C2—H21	107.5	C3—O8—C9	109.8 (2)
O11—C2—H21	110.6	C1—C9—O8	109.9 (2)
C2—C3—C4	115.9 (2)	C1—C9—O10	128.2 (2)
C2—C3—O8	103.3 (2)	O8—C9—O10	121.9 (2)
C4—C3—O8	108.5 (2)	C2—O11—H6	108.3
C2—C3—H31	110.5	C1—O12—H4	114.0
C4—C3—H31	108.2	C1—C13—H131	108.7
O8—C3—H31	110.3	C1—C13—H132	106.3
C3—C4—C5	113.5 (2)	H131—C13—H132	110.1
C3—C4—O7	108.5 (2)	C1—C13—H133	112.3
C5—C4—O7	110.6 (2)	H131—C13—H133	110.1
C3—C4—H41	108.4	H132—C13—H133	109.2

Hydrogen-bond geometry (Å, °)

D—H···A	D—H	H···A	D···A	D—H···A
O12—H4···O7 ⁱ	0.88	1.84	2.723 (2)	178
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