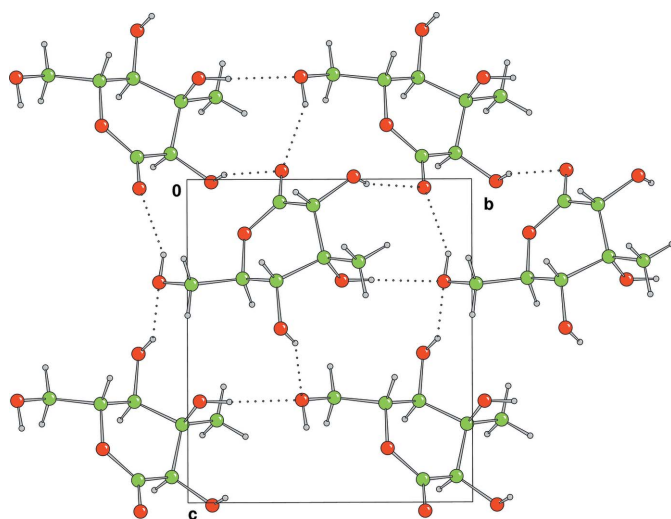


**Figure 1**

The molecular structure of (4), with displacement ellipsoids drawn at the 50% probability level. H atoms are shown as spheres of arbitrary radii.



**Figure 2**

A projection along the *a* axis of part of a hydrogen-bonded sheet of (I). There are no strong interactions between the sheets. Hydrogen bonds are shown as dotted lines.

The value of the Kiliani reaction on 2-*C*-carbon-substituted carbohydrates in the synthesis of 3-*C*-hydroxymethyl branched sugars (Parker *et al.*, 2006; Simone *et al.*, 2007) and 3-*C*-methyl branched sugars (Bream *et al.*, 2006) has been established. Under completely environmentally friendly aqueous conditions, the reaction of cyanide in water with 2-*C*-methyl-*D*-ribose, (2), derived from 2-*C*-methyl-*D*-ribonolactone, (1) (Hotchkiss *et al.*, 2006), gave a major product which crystallized from the reaction mixture. X-ray crystallographic analysis shows (Fig. 1) that the structure is the title compound, (4), removing ambiguities as to the stereochemistry at the new *C*-2 chiral centre and the ring size of the lactone. The minor product is most likely the five-membered ring altrono-lactone, (3). The strain in five-membered ring lactones is generally considerably less than in six-membered ring lactones (Luisa *et al.*, 1990; Brown *et al.*, 1989). Compound (4) is thus a very rare example of the preferential formation of a six-membered ring lactone. Its *C*-2 isomer crystallizes as the 2-*C*-methyl-*D*-allono-

1,4-lactone, (5), rather than the six-membered ring isomer, (6) (Harding *et al.*, 2005). The absolute configuration of compound (4) was determined by the use of the *D*-sugar (1) as the starting material.

The isolated molecule of (4) (Fig. 1) shows no unusual bond lengths or angles, in spite of the strain mentioned above. The largest differences from the *MOGUL* norms (Bruno *et al.*, 2004) are *C*6—*O*7 (00.01 Å; *MOGUL* s.u. 0.02 Å) and *C*3—*C*5—*O*4 (5.0°; *MOGUL* s.u. 2.2°).

The crystal structure of (4) is composed of hydrogen-bonded sheets of molecules lying parallel to the *bc* plane. Both the ketonic atom *O*8 and the hydroxyl atom *O*10 act as acceptors for two hydrogen bonds (Table 1 and Fig. 2).

## Experimental

3-*C*-Methyl-*D*-allono-1,5-lactone, (4), was crystallized from a 3:1:1 mixture of ethyl acetate, methanol and cyclohexane. Analysis: m.p. 421–423 K;  $[\alpha]_{\text{D}}^{23}$  63.5 (*c*, 0.795 in MeOH).

### Crystal data

$\text{C}_7\text{H}_{12}\text{O}_6$	$Z = 2$
$M_r = 192.17$	$D_x = 1.551 \text{ Mg m}^{-3}$
Monoclinic, $P2_1$	Mo $K\alpha$ radiation
$a = 5.6603(2) \text{ \AA}$	$\mu = 0.14 \text{ mm}^{-1}$
$b = 8.0045(2) \text{ \AA}$	$T = 150 \text{ K}$
$c = 9.3242(3) \text{ \AA}$	Needle, colourless
$\beta = 103.1470(13)^\circ$	$0.20 \times 0.05 \times 0.05 \text{ mm}$
$V = 411.39(2) \text{ \AA}^3$	

### Data collection

Nonius KappaCCD area-detector diffractometer	5511 measured reflections
$\omega$ scans	1000 independent reflections
Absorption correction: multi-scan ( <i>DENZO</i> and <i>SCALEPACK</i> ; Otwinowski & Minor, 1997)	930 reflections with $I > 2\sigma(I)$
$T_{\text{min}} = 0.75$ , $T_{\text{max}} = 1.0$	$R_{\text{int}} = 0.050$
	$\theta_{\text{max}} = 27.4^\circ$

### Refinement

Refinement on $F^2$	$w = 1/[\sigma^2(F^2) + (0.02P)^2 + 0.12P]$
$R[F^2 > 2\sigma(F^2)] = 0.031$	where $P = [\max(F_o^2, 0) + 2F_c^2]/3$
$wR(F^2) = 0.076$	$(\Delta/\sigma)_{\text{max}} < 0.001$
$S = 1.16$	$\Delta\rho_{\text{max}} = 0.29 \text{ e \AA}^{-3}$
1000 reflections	$\Delta\rho_{\text{min}} = -0.26 \text{ e \AA}^{-3}$
118 parameters	
H-atom parameters constrained	

**Table 1**

Hydrogen-bond geometry (Å, °).

<i>D</i> — <i>H</i> ... <i>A</i>	<i>D</i> — <i>H</i>	<i>H</i> ... <i>A</i>	<i>D</i> ... <i>A</i>	<i>D</i> — <i>H</i> ... <i>A</i>
<i>O</i> 10— <i>H</i> 7... <i>O</i> 8 <sup>i</sup>	0.84	2.08	2.856 (2)	153
<i>O</i> 7— <i>H</i> 8... <i>O</i> 8 <sup>ii</sup>	0.85	2.06	2.832 (2)	152
<i>O</i> 11— <i>H</i> 12... <i>O</i> 10 <sup>iii</sup>	0.82	2.09	2.836 (2)	152
<i>O</i> 12— <i>H</i> 3... <i>O</i> 10 <sup>iv</sup>	0.83	2.07	2.901 (2)	173

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

In the absence of significant anomalous scattering, Friedel pairs were merged and the absolute configuration assigned from the starting material.

The relatively large ratio of minimum to maximum corrections applied in the multiscan process (1:1.33) reflects changes in the illuminated volume of the very thin needle-like crystal. These changes were kept to a minimum and were taken into account (Görbitz, 1999) by multiscan interframe scaling (*DENZO* and *SCALEPACK*; Otwinowski & Minor, 1997).

The H atoms were all located in a difference map, but 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, with C–H in the range 0.93–0.98 Å and O–H = 0.82 Å, and with  $U_{\text{iso}}(\text{H}) = 1.2\text{--}1.5U_{\text{eq}}(\text{parent atom})$ , after which the positions were refined with riding constraints.

Data collection: *COLLECT* (Nonius, 1997–2001); cell refinement: *DENZO* and *SCALEPACK* (Otwinowski & Minor, 1997); data reduction: *DENZO* and *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*.

The generous gift of 2-*C*-methyl-*D*-ribonolactone from Novartis Pharma AG, Basel, is gratefully acknowledged.

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