

1-Deoxy-D-galactitol (L-fucitol)

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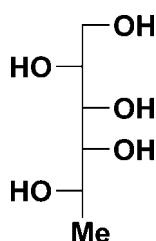
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Key indicators: single-crystal X-ray study; $T = 150\text{ K}$; mean $\sigma(\text{C}-\text{C}) = 0.004\text{ \AA}$; R factor = 0.040; wR factor = 0.111; data-to-parameter ratio = 10.0.

1-Deoxy-D-galactitol, $\text{C}_6\text{H}_{14}\text{O}_5$, exists in the crystalline form as hydrogen-bonded layers of molecules running parallel to the ac plane, with each molecule acting as a donor and acceptor of five hydrogen bonds.

Related literature

For related literature, see: Yoshihara *et al.* (2008); Jones *et al.* (2007); Görbitz (1999); Izumori (2002, 2006); Prince (1982); Watkin (1994).

**Experimental***Crystal data*

$\text{C}_6\text{H}_{14}\text{O}_5$
 $M_r = 166.17$
Monoclinic, $P2_1$
 $a = 4.8486 (3)\text{ \AA}$
 $b = 4.8827 (3)\text{ \AA}$
 $c = 16.8354 (13)\text{ \AA}$
 $\beta = 92.856 (2)^\circ$

$V = 398.07 (5)\text{ \AA}^3$
 $Z = 2$
Mo $K\alpha$ radiation
 $\mu = 0.12\text{ mm}^{-1}$
 $T = 150\text{ K}$
 $0.15 \times 0.15 \times 0.05\text{ mm}$

Data collection

Nonius KappaCCD diffractometer
Absorption correction: multi-scan
(*DENZO/SCALEPACK*;
Otwinowski & Minor, 1997)
 $T_{\min} = 0.81$, $T_{\max} = 0.99$
2786 measured reflections
998 independent reflections
804 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.038$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.040$
 $wR(F^2) = 0.111$
 $S = 0.88$
998 reflections
100 parameters
1 restraint
H-atom parameters constrained
 $\Delta\rho_{\max} = 0.34\text{ e \AA}^{-3}$
 $\Delta\rho_{\min} = -0.31\text{ e \AA}^{-3}$

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

$D-\text{H} \cdots A$	$D-\text{H}$	$\text{H} \cdots A$	$D \cdots A$	$D-\text{H} \cdots A$
O4—H1 \cdots O6 ⁱ	0.83	1.91	2.691 (4)	155
O9—H3 \cdots O4 ⁱⁱ	0.83	1.97	2.753 (4)	156
O6—H4 \cdots O1 ⁱⁱⁱ	0.81	2.10	2.758 (4)	138
O1—H9 \cdots O9 ^{iv}	0.85	1.85	2.684 (4)	166
O11—H10 \cdots O11 ^v	0.84	2.01	2.828 (4)	163

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

Data collection: *COLLECT* (Nonius, 2001); 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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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: LH2653).

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supporting information

Acta Cryst. (2008). E64, o1429 [doi:10.1107/S1600536808020345]

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S1. Comment

The methodology developed by Izumori (2002, 2006) for the interconversion of tetroses, pentoses and hexoses by enzymatic oxidation, inversion at C3 with a single epimerase, and reduction to the aldose has been seen to be generally applicable for the 1-deoxy ketohexoses (Yoshihara *et al.*, 2008). This methodology could allow access to rare monosaccharides in water in large amounts. An example of this is the subsequent formation of 1-deoxy-D-galactitol **2** by hydrogenation of L-fucose **1** (Fig. 1) which subsequently could be oxidized enzymatically to 1-deoxy-D-tagatose (Jones *et al.*, 2007) **3**.

If the terminal hydroxyl group and H atoms are ignored there is a pseudo centre of symmetry between C2 and C3 (Fig. 2). The crystal structure exists of hydrogen-bonded layers of molecules running parallel to the *c*-axis (Fig. 3). Each molecule acts as a donor and acceptor of 5 hydrogen bonds, all intra-molecular hydrogen bonds have been omitted.

S2. Experimental

The title compound was recrystallized from methanol: m.p. 420–422 K; $[\alpha]_D^{21} +1.6$ (*c*, 1.13 in H₂O) [Lit. (Yoshihara *et al.*, 2008) for enantiomer $[\alpha]_D^{20} -1.9$ (*c*, 1.0 in H₂O)].

S3. Refinement

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

The relatively large ratio of minimum to maximum corrections applied in the multiscan process (1:1.22 reflect changes in the illuminated volume of the crystal. Changes in illuminated volume were kept to a minimum, and were taken into account (Görbitz, 1999) by the multi-scan inter-frame scaling (*DENZO/SCALEPACK*, Otwinowski & Minor, 1997).

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

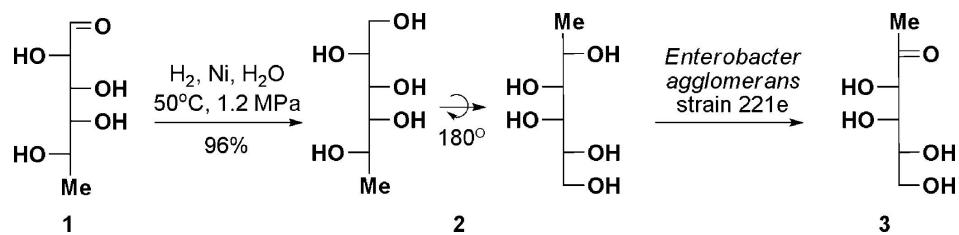


Figure 1

Synthetic scheme.

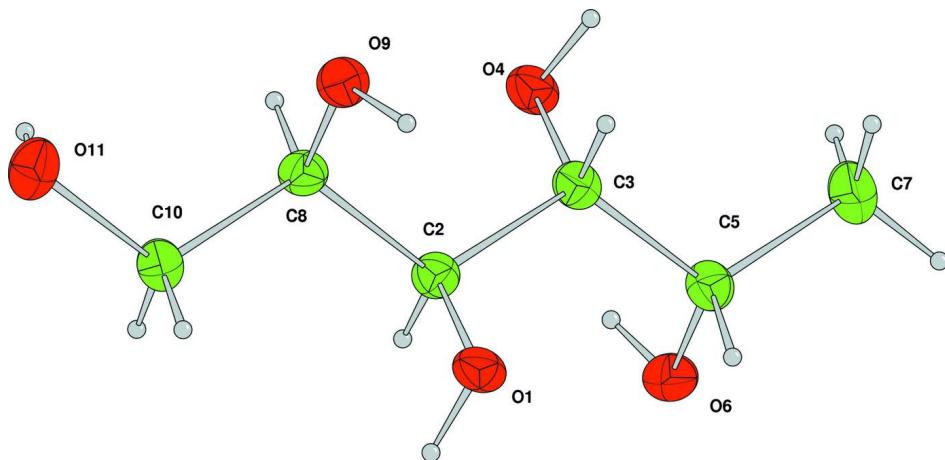
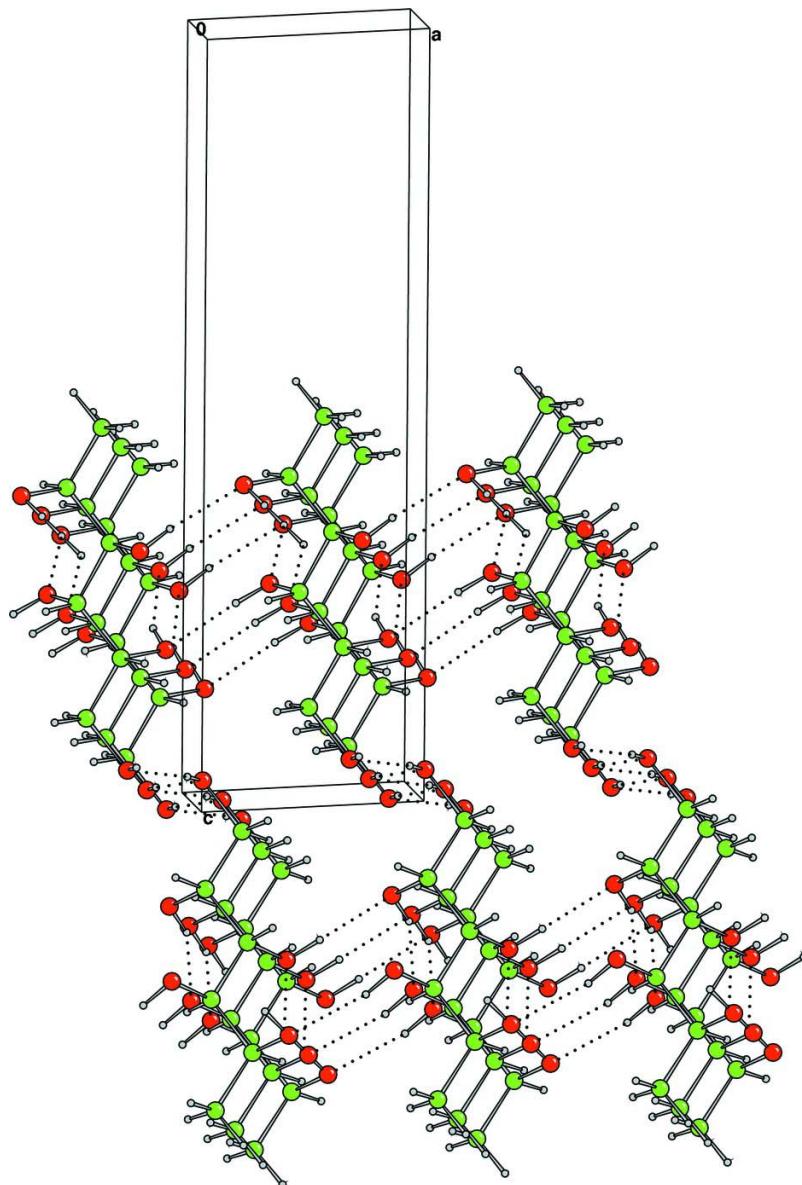


Figure 2

The title compound with displacement ellipsoids drawn at the 50% probability level. H atoms are shown as spheres of arbitrary radius.

**Figure 3**

The packing diagram for the title compound projected along the *b*-axis. Hydrogen bonds are shown as dotted lines.

1-Deoxy-D-galactitol

Crystal data

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 $M_r = 166.17$
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 $\beta = 92.856 (2)^\circ$
 $V = 398.07 (5) \text{ \AA}^3$
 $Z = 2$

$F(000) = 180$
 $D_x = 1.386 \text{ Mg m}^{-3}$
Mo $K\alpha$ radiation, $\lambda = 0.71073 \text{ \AA}$
Cell parameters from 844 reflections
 $\theta = 5\text{--}27^\circ$
 $\mu = 0.12 \text{ mm}^{-1}$
 $T = 150 \text{ K}$
Block, colourless
 $0.15 \times 0.15 \times 0.05 \text{ mm}$

Data collection

Nonius KappaCCD
diffractometer
Graphite monochromator
 ω scans
Absorption correction: multi-scan
(DENZO/SCALEPACK; Otwinowski & Minor,
1997)
 $T_{\min} = 0.81$, $T_{\max} = 0.99$

2786 measured reflections
998 independent reflections
804 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.038$
 $\theta_{\max} = 27.4^\circ$, $\theta_{\min} = 5.4^\circ$
 $h = -6 \rightarrow 6$
 $k = -5 \rightarrow 6$
 $l = -21 \rightarrow 21$

Refinement

Refinement on F^2
Least-squares matrix: full
 $R[F^2 > 2\sigma(F^2)] = 0.040$
 $wR(F^2) = 0.111$
 $S = 0.88$
998 reflections
100 parameters
1 restraint
Primary atom site location: structure-invariant
direct methods
Hydrogen site location: inferred from
neighbouring sites

H-atom parameters constrained
Method, part 1, Chebychev polynomial,
(Watkin, 1994; Prince, 1982) [weight] =
 $1.0/[A_0*T_0(x) + A_1*T_1(x) \dots + A_{n-1}*T_{n-1}(x)]$
where A_i are the Chebychev coefficients listed
below and $x = F/F_{\max}$ Method = Robust
Weighting (Prince, 1982) $W = [\text{weight}] *$
 $[1-(\Delta F/6*\sigma F)^2]^2$ A_i are: 17.0 25.0 12.0
3.16
 $(\Delta/\sigma)_{\max} = 0.000240$
 $\Delta\rho_{\max} = 0.34 \text{ e } \text{\AA}^{-3}$
 $\Delta\rho_{\min} = -0.31 \text{ e } \text{\AA}^{-3}$

Fractional atomic coordinates and isotropic or equivalent isotropic displacement parameters (\AA^2)

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$
O1	0.4779 (4)	0.0226 (5)	0.76245 (11)	0.0217
C2	0.6328 (6)	0.2631 (7)	0.78168 (17)	0.0186
C3	0.7866 (6)	0.3389 (7)	0.70769 (17)	0.0189
O4	0.9430 (4)	0.5805 (5)	0.72728 (12)	0.0227
C5	0.5946 (6)	0.3936 (7)	0.63490 (17)	0.0207
O6	0.4117 (4)	0.6179 (5)	0.64879 (12)	0.0238
C7	0.7550 (7)	0.4471 (9)	0.56067 (18)	0.0330
C8	0.8283 (6)	0.2108 (7)	0.85426 (17)	0.0190
O9	1.0094 (4)	-0.0141 (5)	0.84026 (12)	0.0222
C10	0.6698 (6)	0.1572 (7)	0.92859 (17)	0.0236
O11	0.8526 (4)	0.1176 (5)	0.99759 (12)	0.0260
H21	0.5071	0.4100	0.7945	0.0249*
H31	0.9082	0.1875	0.6971	0.0263*
H51	0.4763	0.2307	0.6253	0.0282*
H71	0.6272	0.4510	0.5138	0.0515*
H72	0.8900	0.3047	0.5550	0.0518*
H73	0.8493	0.6223	0.5674	0.0506*
H81	0.9485	0.3709	0.8670	0.0243*
H101	0.5642	-0.0123	0.9193	0.0325*
H102	0.5415	0.3107	0.9363	0.0333*
H1	1.0737	0.5438	0.6989	0.0372*
H3	0.9415	-0.1296	0.8087	0.0364*
H4	0.5121	0.7060	0.6789	0.0402*
H9	0.3277	0.0397	0.7859	0.0353*

H10	0.9076	0.2813	0.9992	0.0410*
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Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
O1	0.0179 (9)	0.0228 (13)	0.0249 (10)	-0.0051 (9)	0.0048 (8)	-0.0061 (10)
C2	0.0180 (13)	0.0189 (15)	0.0189 (13)	-0.0011 (11)	0.0010 (10)	0.0011 (12)
C3	0.0196 (13)	0.0173 (15)	0.0202 (13)	-0.0016 (12)	0.0031 (11)	-0.0029 (12)
O4	0.0212 (10)	0.0235 (13)	0.0237 (9)	-0.0059 (10)	0.0057 (8)	-0.0040 (10)
C5	0.0210 (14)	0.0218 (17)	0.0196 (13)	0.0007 (13)	0.0029 (11)	-0.0017 (12)
O6	0.0188 (9)	0.0271 (13)	0.0254 (10)	0.0014 (10)	0.0003 (8)	-0.0008 (11)
C7	0.0320 (17)	0.048 (2)	0.0192 (14)	0.0027 (17)	0.0047 (12)	0.0033 (16)
C8	0.0166 (13)	0.0198 (15)	0.0204 (13)	0.0021 (12)	0.0006 (10)	-0.0004 (12)
O9	0.0206 (10)	0.0227 (12)	0.0233 (10)	0.0015 (10)	0.0011 (8)	-0.0047 (10)
C10	0.0223 (14)	0.031 (2)	0.0179 (13)	0.0020 (13)	0.0023 (11)	-0.0001 (13)
O11	0.0323 (11)	0.0248 (11)	0.0206 (9)	-0.0028 (11)	-0.0024 (8)	0.0022 (10)

Geometric parameters (\AA , $^\circ$)

O1—C2	1.423 (4)	O6—H4	0.809
O1—H9	0.849	C7—H71	0.979
C2—C3	1.529 (4)	C7—H72	0.963
C2—C8	1.530 (4)	C7—H73	0.974
C2—H21	0.972	C8—O9	1.433 (4)
C3—O4	1.432 (4)	C8—C10	1.523 (4)
C3—C5	1.525 (4)	C8—H81	0.992
C3—H31	0.968	O9—H3	0.832
O4—H1	0.832	C10—O11	1.439 (4)
C5—O6	1.436 (4)	C10—H101	0.982
C5—C7	1.527 (4)	C10—H102	0.987
C5—H51	0.989	O11—H10	0.843
C2—O1—H9	105.6	C5—C7—H71	109.6
O1—C2—C3	106.7 (2)	C5—C7—H72	109.6
O1—C2—C8	110.0 (3)	H71—C7—H72	109.9
C3—C2—C8	112.6 (2)	C5—C7—H73	108.1
O1—C2—H21	109.2	H71—C7—H73	110.5
C3—C2—H21	109.8	H72—C7—H73	109.1
C8—C2—H21	108.5	C2—C8—O9	110.9 (2)
C2—C3—O4	106.6 (2)	C2—C8—C10	111.5 (2)
C2—C3—C5	113.2 (2)	O9—C8—C10	110.0 (3)
O4—C3—C5	109.6 (3)	C2—C8—H81	112.0
C2—C3—H31	106.9	O9—C8—H81	106.3
O4—C3—H31	110.6	C10—C8—H81	105.8
C5—C3—H31	109.8	C8—O9—H3	113.5
C3—O4—H1	95.8	C8—C10—O11	111.8 (2)
C3—C5—O6	111.1 (2)	C8—C10—H101	107.1
C3—C5—C7	111.9 (2)	O11—C10—H101	108.1

O6—C5—C7	110.3 (3)	C8—C10—H102	108.9
C3—C5—H51	108.5	O11—C10—H102	111.3
O6—C5—H51	106.4	H101—C10—H102	109.5
C7—C5—H51	108.6	C10—O11—H10	94.6
C5—O6—H4	98.7		

Hydrogen-bond geometry (Å, °)

D—H···A	D—H	H···A	D···A	D—H···A
O4—H1···O6 ⁱ	0.83	1.91	2.691 (4)	155
O9—H3···O4 ⁱⁱ	0.83	1.97	2.753 (4)	156
O6—H4···O1 ⁱⁱⁱ	0.81	2.10	2.758 (4)	138
O6—H4···O4	0.81	2.29	2.842 (4)	126
O1—H9···O9 ^{iv}	0.85	1.85	2.684 (4)	166
O11—H10···O11 ^v	0.84	2.01	2.828 (4)	163

Symmetry codes: (i) $x+1, y, z$; (ii) $x, y-1, z$; (iii) $x, y+1, z$; (iv) $x-1, y, z$; (v) $-x+2, y+1/2, -z+2$.