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(-)-*epi*-Inosose-2¹

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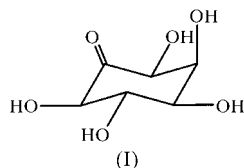
Data validation number: IUC0000309

The structure of the title compound, C₆H₁₀O₆, was determined to confirm the position of the keto group in the molecule prepared enantioselectively by a bioconversion from *myo*-inositol. There are two independent molecules showing similar geometry.

Comment

myo-Inositol has only one axial hydroxyl group and has essentially mirror symmetry. Crystal structures have been reported for *myo*-inositol (Rabinowitz & Kraut, 1964), *myo*-inositol dihydrate (Lomer *et al.*, 1963), *myo*-inositol tetraaquamagnesium dichloride (Blank, 1973) and *myo*-inositol calcium bromide pentahydrate (Cook & Bugg, 1973). The crystal structure and molecular dynamics analysis of an optically active *myo*-inositol derivative have been reported recently (Dillen *et al.*, 2000).

The title compound, (I), was prepared enantioselectively by a bioconversion from *myo*-inositol (Takahashi *et al.*, 2000).



Experimental

Crystals of (I) were grown from an aqueous ethanol solution.

¹ Systematic name: (-)-2L-2,3,4,6/5-pentahydroxycyclohexanone.

Crystal data

C₆H₁₀O₆
M_r = 178.14
Monoclinic, P2₁
a = 11.197 (2) Å
b = 8.932 (2) Å
c = 6.976 (2) Å
β = 90.21 (2)°
V = 697.7 (3) Å³
Z = 4

D_x = 1.696 Mg m⁻³
Mo Kα radiation
Cell parameters from 25 reflections
θ = 13.9–15.0°
μ = 0.155 mm⁻¹
T = 298 (1) K
Prism, colourless
0.7 × 0.4 × 0.15 mm

Data collection

Rigaku AFC-7R diffractometer
θ–2θ scans
2250 measured reflections
2151 independent reflections
2086 reflections with I > 2σ(I)
R_{int} = 0.005
θ_{max} = 30°

h = 0 → 16
k = 0 → 13
l = -10 → 10
3 standard reflections every 150 reflections
intensity decay: none

Refinement

Refinement on F²
R(F) = 0.024
wR(F²) = 0.068
S = 1.13
2151 reflections
298 parameters
All H-atom parameters refined

w = 1/[σ²(F_o²) + (0.0417P)² + 0.0965P]
where P = (F_o² + 2F_c²)/3
(Δ/σ)_{max} < 0.001
Δρ_{max} = 0.30 e Å⁻³
Δρ_{min} = -0.16 e Å⁻³
Extinction correction: SHELXL97 (Sheldrick, 1997)
Extinction coefficient: 0.027 (5)

Table 1

Selected geometric parameters (Å).

O1—C7	1.205 (2)	O51—C57	1.207 (2)
O2—C8	1.419 (2)	O52—C58	1.401 (2)
O3—C9	1.425 (2)	O53—C59	1.440 (2)
O4—C10	1.428 (2)	O54—C60	1.427 (2)
O5—C11	1.424 (2)	O55—C61	1.424 (2)
O6—C12	1.418 (2)	O56—C62	1.414 (2)

Table 2

Hydrogen-bonding geometry (Å, °).

D—H...A	D—H	H...A	D...A	D—H...A
O2—H2...O53 ⁱ	0.76 (3)	2.09 (3)	2.856 (2)	177 (3)
O3—H3...O54 ⁱ	0.84 (3)	1.96 (3)	2.786 (2)	166 (3)
O4—H4...O53 ⁱⁱ	0.73 (3)	2.49 (3)	3.123 (2)	145 (3)
O5—H5...O56 ⁱⁱⁱ	0.82 (3)	1.97 (3)	2.785 (2)	173 (4)
O6—H6...O4 ^{iv}	0.76 (3)	2.10 (3)	2.828 (2)	163 (3)
O52—H52...O3 ^v	0.85 (4)	2.04 (4)	2.856 (2)	161 (3)
O53—H53...O51 ^v	0.81 (3)	2.06 (3)	2.845 (2)	164 (3)
O54—H54...O55 ^{vi}	0.83 (3)	1.94 (3)	2.767 (2)	172 (2)
O55—H55...O6 ^{vii}	0.81 (3)	1.98 (3)	2.782 (2)	170 (3)
O56—H56...O2	0.89 (4)	2.06 (4)	2.938 (2)	170 (3)

Symmetry codes: (i) 1 - x, ½ + y, 1 - z; (ii) 1 - x, ½ + y, -z; (iii) 2 - x, ½ + y, -z; (iv) 2 - x, y - ½, -z; (v) 1 - x, y - ½, -z; (vi) 1 - x, y - ½, 1 - z; (vii) 2 - x, y - ½, 1 - z.

All H atoms were located from difference syntheses and refined isotropically; the O—H and C—H lengths are 0.73 (3)–0.85 (4) and 0.91 (2)–1.01 (2) Å, respectively. The absolute structure was assumed based on the known absolute configuration of the title compound (Posternak, 1946).

Data collection: MSC/AFC Diffractometer Control Software (Molecular Structure Corporation, 1993); cell refinement: MSC/AFC Diffractometer Control Software; data reduction: TEXSAN (Molecular Structure Corporation, 1999); program(s) used to solve struc-

ture: *SIR92* (Altomare *et al.*, 1994); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); software used to prepare material for publication: *TEXSAN*.

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