

(2*R*,3*R*,4*R*,5*S*)-3,4,5-Trihydroxy-2-(2-hydroxyethyl)piperidinium chloride

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Received 20 December 1999

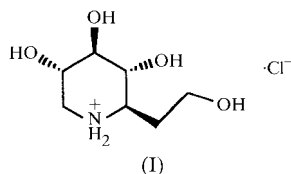
Accepted 1 March 2000

Data validation number: IUC0000050

The absolute configuration at the new stereogenic centre during the key step of the total synthesis was established by X-ray analysis of the title compound, C₇H₁₅NO₄⁺·Cl⁻.

Comment

The title compound [(I); alternative name: 1,5,6-trideoxy-1,5-imino-D-*gluco*-heptitol hydrochloride] was prepared by palladium(II)-catalyzed aminocarbonylation of (2*S*,3*S*,4*R*)-1-benzylamino-2,3-di-*O*-benzylhex-5-ene-2,3,4-triol with subsequent lactone-ring opening and deprotection (Szolcsányi *et al.*, 2000).



The bond lengths are consistent with average values in the usual sources [*International Tables for Crystallography* (1992, Vol. C, Table 9.5.1.1)]. The Cl⁻ anion is bonded in the structure by hydrogen bonds [Cl⁻··H15 2.225 Å, Cl⁻··H16 2.725 Å, Clⁱⁱⁱ··H12 2.420 Å and Clⁱⁱⁱ··H1 2.300 Å; symmetry codes: (i) $-x + 2, y - \frac{1}{2}, -z$; (ii) $-x + 1, y - \frac{1}{2}, -z$; (iii) $x + 1, y, z$].

Experimental

The title compound was prepared by hydrogenolysis of the benzylated diol with H₂ on Pd/C at room temperature in EtOH containing a few drops of aqueous HCl. The crude product was purified by ion-exchange chromatography (H⁺ form, DOWEX 50WX8-400) and recrystallized from methanol/*tert*-butyl methyl ether (m.p. 450–452 K, $[\alpha]_D^{20} = +30, c = 0.55, \text{MeOH}$).

Crystal data

C₇H₁₆NO₄⁺·Cl⁻
M_r = 213.66
Monoclinic, *P*2₁
a = 6.8910 (10) Å
b = 7.3450 (10) Å
c = 9.497 (2) Å
β = 92.46 (3)°
V = 480.24 (14) Å³
Z = 2

D_x = 1.478 Mg m⁻³
Mo Kα radiation
Cell parameters from 20 reflections
θ = 4.53–10.24°
μ = 0.382 mm⁻¹
T = 293 (2) K
Rectangular, colourless
0.6 × 0.5 × 0.2 mm

Data collection

Syntex P2₁ diffractometer
θ–2θ scans
Absorption correction: ψ scan
(North *et al.*, 1968)
T_{min} = 0.791, T_{max} = 0.926
1663 measured reflections
1566 independent reflections
1222 reflections with I > 2σ(I)

R_{int} = 0.048
θ_{max} = 30.08°
h = 0 → 9
k = -10 → 10
l = -13 → 13
2 standard reflections
every 100 reflections
intensity decay: 15%

Refinement

Refinement on F²
R[F² > 2σ(F²)] = 0.056
wR(F²) = 0.157
S = 0.933
1461 reflections
118 parameters
H-atom parameters constrained

w = 1/[σ²(F_o²) + (0.1308P)² + 0.3128P]
where P = (F_o² + 2F_c²)/3
(Δ/σ)_{max} < 0.001
Δρ_{max} = 0.84 e Å⁻³
Δρ_{min} = -0.54 e Å⁻³

Table 1

Selected geometric parameters (Å, °).

O1–C1	1.414 (6)	O4–C6	1.419 (5)
O2–C4	1.412 (4)	N1–C7	1.479 (5)
O3–C5	1.425 (4)	N1–C3	1.500 (5)
C7–N1–C3	114.2 (3)	C5–C4–C3	113.1 (3)
O1–C1–C2	107.6 (3)	O3–C5–C6	108.2 (3)
C1–C2–C3	113.8 (3)	O3–C5–C4	110.8 (3)
N1–C3–C2	109.8 (3)	C6–C5–C4	110.4 (3)
N1–C3–C4	109.6 (3)	O4–C6–C5	110.9 (3)
C2–C3–C4	110.9 (3)	O4–C6–C7	108.6 (3)
O2–C4–C5	110.1 (3)	C5–C6–C7	108.8 (3)
O2–C4–C3	106.0 (3)	N1–C7–C6	109.3 (3)

The number of Friedel pairs in the data set is 60.

Data collection: *Syntex P2₁ Software* (Syntex, 1973); cell refinement: *Syntex P2₁ Software*; data reduction: *XP21* (Pavelčík, 1993); program(s) used to solve structure: *SHELXS86* (Sheldrick, 1990); program(s) used to refine structure: *SHELXL93* (Sheldrick, 1993); software used to prepare material for publication: *SHELXL93*.

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