Acta Crystallographica Section E
Structure Reports Online
ISSN 1600-5368

N-(4-Chlorophenyl)-2-deoxy-α-L-ribo-pyranosylamine

Pei-Hua Shang,[a] Chang-Mei Cheng,[a]* Yong-Chong Yang,[a] Ru-Ji Wang[b] and Yu-Fen Zhao[a]

[a]Key Laboratory for Bioorganic Phosphorus Chemistry and Chemical Biology, Ministry of Education, Department of Chemistry, Tsinghua University, Beijing 100084, People's Republic of China, and [b]Department of Chemistry, Tsinghua University, Beijing 100084, People's Republic of China
Correspondence e-mail: shangpeihua05@mail.tsinghua.org.cn

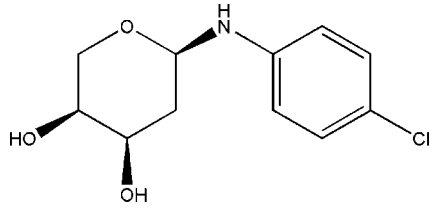
Received 19 March 2008; accepted 23 April 2008

Key indicators: single-crystal X-ray study; T = 295 K; mean σ(C–C) = 0.004 Å; R factor = 0.037; wR factor = 0.084; data-to-parameter ratio = 14.8.

In the crystal structure of the title compound, $C_{11}H_{14}ClNO_3$, intermolecular hydrogen bonds link molecules in the *ab* plane, forming layers that stack along the *c* axis.

Related literature

For related literature, see: Durette *et al.* (1978); Ganem (1966); Katzen (1979); Bridiau *et al.* (2007); Ojala *et al.* (2000).



Experimental

Crystal data

$C_{11}H_{14}ClNO_3$
M_r = 243.68
Orthorhombic, $P2_12_12_1$
a = 6.5305 (8) Å
b = 7.9857 (9) Å
c = 22.496 (3) Å
V = 1173.2 (3) Å^3
Z = 4
Mo $K\alpha$ radiation
μ = 0.32 mm^{-1}
T = 295 (2) K
0.4 × 0.2 × 0.1 mm

Data collection

Bruker P4 diffractometer
Absorption correction: none
2581 measured reflections
2172 independent reflections
1690 reflections with $I > 2\sigma(I)$
R_{int} = 0.025
3 standard reflections
every 97 reflections
intensity decay: none

Refinement

$R[F^2 > 2\sigma(F^2)]$ = 0.037
$wR(F^2)$ = 0.084
S = 1.03
2172 reflections
147 parameters
H-atom parameters constrained
$\Delta\rho_{max}$ = 0.15 e Å^{-3}
$\Delta\rho_{min}$ = −0.17 e Å^{-3}
Absolute structure: Flack (1983), 880 Friedel pairs
Flack parameter: 0.09 (11)

Table 1
Hydrogen-bond geometry (Å, °).

D—H···A	D—H	H···A	D···A	D—H···A
O2—H2*C*···O3^{i}	0.82	1.93	2.739 (2)	170
O3—H3*B*···O1^{i}	0.82	1.98	2.797 (2)	175
N1—H1*B*···O2^{ii}	0.92	2.08	2.994 (3)	173

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

Data collection: *XSCANS* (Bruker, 1997); cell refinement: *XSCANS*; data reduction: *XSCANS*; program(s) used to solve structure: *SHELXTL* (Sheldrick, 2008); program(s) used to refine structure: *SHELXTL*; molecular graphics: *SHELXTL*; software used to prepare material for publication: *SHELXTL*.

We are grateful to the National Science Foundation of China (project grant No. 20132020).

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: PK2090).

References

Bridiau, N., Benmansour, M., Legoy, M. D. & Maugard, T. (2007). *Tetrahedron*, **63**. 4178–4183.
Bruker (1997). *XSCANS*. Bruker AXS Inc.,Madison, Wisconsin, USA.
Durette, P. L., Bugianesi, R. L., Ponpipom, M. M., Shen, T. Y., Cascieri, M. A., Glitzer, M. S. & Katzen, H. M. (1978). *J. Med. Chem.* **21**, 854–859.
Flack, H. D. (1983). *Acta Cryst.* A**39**, 876–881.
Ganem, B. (1966). *Acc. Chem. Res.* **29**, 340–347.
Katzen, H. M. (1979). *J. Biol. Chem.* **254**, 2983–2992.
Ojala, W. H., Oatman, J. M. & Ojala, C. R. (2000). *Carbohydr. Res.* **326**, 104–112.
Sheldrick, G. M. (2008). *Acta Cryst.* A**64**, 112–122.

supporting information

Acta Cryst. (2008). E**64**, o1317 [doi:10.1107/S1600536808011616]

N-(4-Chlorophenyl)-2-deoxy-*α*-L-ribopyranosylamine

Pei-Hua Shang, Chang-Mei Cheng, Yong-Chong Yang, Ru-Ji Wang and Yu-Fen Zhao

S1. Comment

N-Alkyl and *N*-aryl glycosylamines have a wide range of biological activities (Katzen *et al.*, 1979; Ganem, 1966), including insulin-like activity (Durette *et al.*, 1978). They are important as junctures in glycoproteins (Ojala *et al.*, 2000). Glycosylamines can exist either in cyclic or acyclic forms depending on reaction conditions and the particular amine used. Stereo-selective syntheses of *N*-aryl-glycosylamines are uncommon, but a one-pot stereoselective synthesis of beta-*N*-aryl-glycosides in aqueous buffers with purification by semi-preparative HPLC has been reported (Nicolas *et al.*, 2007).

Recently, we found that 4-chlorobenzenamine reacted with 2-deoxy-*L*-ribose in methanol and water to give *N*-*p*-chlorophenyl-2-deoxy-*α*-*L*-ribopyranosylamine as the sole product. Herein we report the synthesis and structure (Fig. 1) of *N*-*p*-chlorophenyl-2-deoxy-*α*-*L*-ribopyranosylamine.

S2. Experimental

The title compound was synthesized by the reaction of 4-chlorobenzenamine with 2-deoxy-*L*-ribose in a mixture of methanol and water. 4-chlorobenzenamine (0.93 g, 10 mmol) in a little methanol was added to a solution of 2-deoxy-*L*-ribose (1.34 g, 10 mmol) in 20 ml water, the solution was stirred at room temperature overnight. A white solid obtained by filtration was washed with ice water, then cold ether, and was dried under pressure. The solid was *N*-*p*-chlorophenyl-2-deoxy-*α*-*L*-ribopyranosylamine (yield: 70%). ^{1}H NMR (300 MHz, DMSO-d$_6$): δ 1.68 (m, 1H), 1.78 (m, 1H), 3.37 (d, 1H), 3.49 (s, 1H), 3.62 (q, 1H), 3.68 (m, 1H), 4.36 (d, 1H), 4.56 (m, 1H), 4.69 (d, 1H), 6.16 (d, 1H), 6.53 (d, 2H), 6.886 (d, 2H). ^{13}C NMR (300 MHz, DMSO-d$_6$): δ 144.2, 129.2, 125.3, 113.4, 80.3, 68.0, 66.8, 65.7, 34.7, 20.1.

S3. Refinement

H atoms were placed in calculated positions with constrained distances of 0.98 Å (R_3CH), 0.97 Å (R_2CH_2), 0.93 Å (R_2CH), 0.82 Å (OH) and 0.9195 Å (NH). U_{iso}(H) values were set to 1.2U_{eq} of the attached atom.

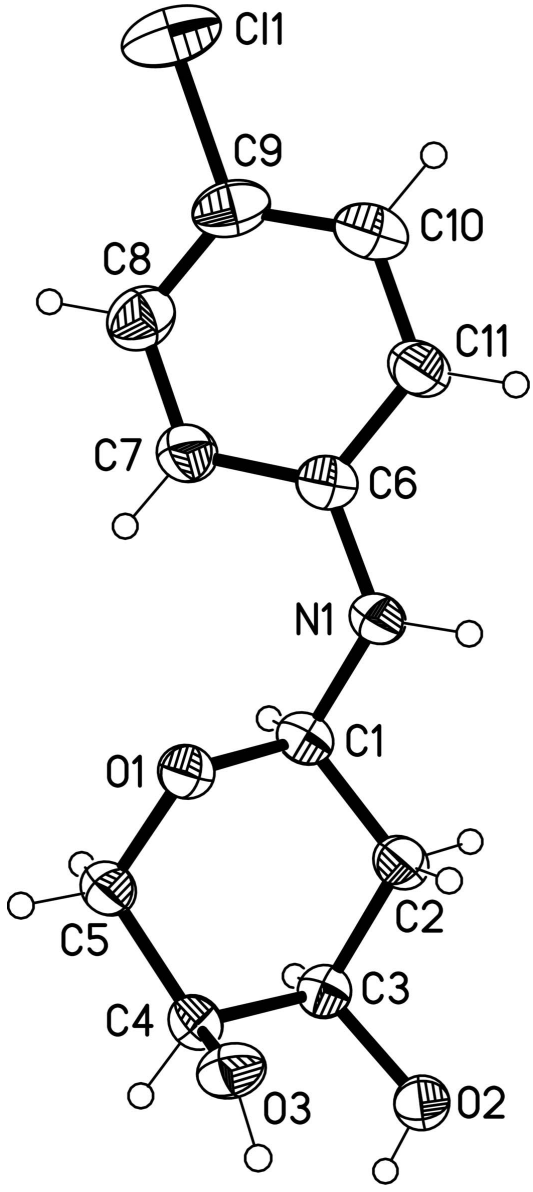


Figure 1

View of the title compound, with displacement ellipsoids drawn at the 35% probability level.

N-(4-Chlorophenyl)-2-deoxy-α-L-ribopyranosylamine

Crystal data

$C_{11}H_{14}ClNO_3$
$M_r = 243.68$
Orthorhombic, $P2_12_12_1$
Hall symbol: P 2ac 2ab
$a = 6.5305\ (8)$ Å
$b = 7.9857\ (9)$ Å
$c = 22.496\ (3)$ Å
$V = 1173.2\ (3)$ Å^3
$Z = 4$
$F(000) = 512$
$D_x = 1.380$ Mg m^{-3}
Mo $K\alpha$ radiation, $\lambda = 0.71073$ Å
Cell parameters from 37 reflections
$\theta = 4.9$–$12.5°$
$\mu = 0.32$ mm^{-1}
$T = 295$ K
Prism, colorless
0.4 × 0.2 × 0.1 mm

Data collection

Bruker P4 diffractometer
Radiation source: fine-focus sealed tube
Graphite monochromator
ω scans
2581 measured reflections
2172 independent reflections
1690 reflections with $I > 2\sigma(I)$

$R_{int} = 0.026$
$\theta_{max} = 25.5°$, $\theta_{min} = 2.7°$
$h = -7 \rightarrow 7$
$k = -9 \rightarrow 9$
$l = -27 \rightarrow 27$
3 standard reflections every 97 reflections
intensity decay: none

Refinement

Refinement on F^2
Least-squares matrix: full
$R[F^2 > 2\sigma(F^2)] = 0.037$
$wR(F^2) = 0.084$
$S = 1.03$
2172 reflections
147 parameters
0 restraints
Primary atom site location: structure-invariant direct methods
Secondary atom site location: difference Fourier map

Hydrogen site location: inferred from neighbouring sites
H-atom parameters constrained
$w = 1/[\sigma^2(F_o^2) + (0.005P)^2 + 0.4P]$ where $P = (F_o^2 + 2F_c^2)/3$
$(\Delta/\sigma)_{max} < 0.001$
$\Delta\rho_{max} = 0.15$ e Å^{-3}
$\Delta\rho_{min} = -0.17$ e Å^{-3}
Absolute structure: Flack (1983), 880 Friedel pairs
Absolute structure parameter: 0.09 (11)

Special details

Geometry. All e.s.d.'s (except the e.s.d. in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell e.s.d.'s are taken into account individually in the estimation of e.s.d.'s in distances, angles and torsion angles; correlations between e.s.d.'s in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell e.s.d.'s is used for estimating e.s.d.'s involving l.s. planes.

Refinement. Refinement of F^2 against ALL reflections. The weighted *R*-factor *wR* and goodness of fit *S* are based on F^2, conventional *R*-factors *R* are based on *F*, with *F* set to zero for negative F^2. The threshold expression of $F^2 > \sigma(F^2)$ is used only for calculating *R*-factors(gt) *etc*. and is not relevant to the choice of reflections for refinement. *R*-factors based on F^2 are statistically about twice as large as those based on *F*, and *R*- factors based on ALL data will be even larger.

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

	x	*y*	*z*	U_{iso}*/U_{eq}
Cl1	0.01956 (18)	−0.12260 (10)	0.00675 (4)	0.0884 (3)
O1	0.3998 (2)	0.56562 (19)	0.16726 (7)	0.0425 (4)
O2	0.2117 (3)	1.0211 (2)	0.23361 (8)	0.0482 (5)
H2C	0.2839	1.1049	0.2316	0.058*
O3	0.5280 (3)	0.7894 (2)	0.25924 (7)	0.0477 (4)
H3B	0.5420	0.8705	0.2812	0.057*
N1	0.0604 (3)	0.4754 (3)	0.16049 (9)	0.0495 (6)
H1B	−0.0305	0.4823	0.1916	0.059*
C1	0.1909 (4)	0.6147 (3)	0.15314 (11)	0.0412 (6)
H1A	0.1850	0.6517	0.1116	0.049*
C2	0.1280 (4)	0.7577 (3)	0.19294 (12)	0.0457 (6)
H2A	0.1197	0.7179	0.2336	0.055*
H2B	−0.0071	0.7960	0.1813	0.055*
C3	0.2759 (4)	0.9032 (3)	0.19013 (10)	0.0392 (6)
H3A	0.2653	0.9554	0.1508	0.047*

C4	0.4948 (4)	0.8459 (3)	0.19940 (10)	0.0412 (6)
H4A	0.5884	0.9386	0.1906	0.049*
C5	0.5384 (4)	0.7017 (3)	0.15789 (11)	0.0451 (6)
H5A	0.5270	0.7400	0.1171	0.054*
H5B	0.6776	0.6631	0.1641	0.054*
C6	0.0505 (4)	0.3402 (3)	0.12239 (11)	0.0450 (6)
C7	0.2014 (5)	0.3069 (4)	0.08000 (12)	0.0549 (7)
H7A	0.3118	0.3797	0.0761	0.066*
C8	0.1893 (5)	0.1674 (4)	0.04379 (12)	0.0591 (8)
H8A	0.2906	0.1472	0.0156	0.071*
C9	0.0287 (6)	0.0594 (3)	0.04936 (11)	0.0567 (8)
C10	−0.1245 (5)	0.0904 (4)	0.08970 (12)	0.0580 (8)
H10A	−0.2348	0.0172	0.0928	0.070*
C11	−0.1145 (5)	0.2305 (3)	0.12566 (12)	0.0527 (7)
H11A	−0.2198	0.2518	0.1525	0.063*

Atomic displacement parameters (Å²)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
Cl1	0.1371 (9)	0.0575 (4)	0.0707 (5)	0.0048 (6)	−0.0236 (6)	−0.0194 (4)
O1	0.0405 (9)	0.0404 (9)	0.0464 (9)	−0.0016 (8)	−0.0010 (8)	−0.0021 (8)
O2	0.0471 (11)	0.0378 (10)	0.0595 (10)	−0.0028 (9)	0.0080 (9)	−0.0056 (9)
O3	0.0561 (11)	0.0442 (9)	0.0429 (9)	0.0035 (10)	−0.0106 (9)	−0.0058 (8)
N1	0.0488 (13)	0.0482 (12)	0.0515 (12)	−0.0157 (11)	0.0108 (11)	−0.0106 (11)
C1	0.0396 (13)	0.0424 (13)	0.0417 (13)	−0.0043 (12)	−0.0045 (11)	0.0018 (12)
C2	0.0380 (14)	0.0423 (14)	0.0568 (15)	−0.0025 (11)	−0.0008 (12)	−0.0025 (13)
C3	0.0419 (14)	0.0359 (13)	0.0399 (12)	0.0022 (11)	−0.0002 (11)	0.0037 (11)
C4	0.0392 (14)	0.0423 (13)	0.0422 (12)	−0.0042 (12)	0.0011 (11)	0.0017 (10)
C5	0.0415 (14)	0.0480 (13)	0.0457 (13)	−0.0077 (13)	0.0025 (12)	−0.0025 (12)
C6	0.0501 (15)	0.0422 (13)	0.0427 (13)	−0.0043 (13)	−0.0031 (12)	0.0007 (11)
C7	0.0566 (17)	0.0569 (17)	0.0513 (15)	−0.0107 (16)	0.0078 (15)	−0.0060 (14)
C8	0.073 (2)	0.0582 (18)	0.0464 (15)	0.0010 (18)	0.0044 (15)	−0.0038 (14)
C9	0.085 (2)	0.0446 (14)	0.0408 (13)	0.0014 (17)	−0.0127 (16)	−0.0011 (12)
C10	0.0685 (19)	0.0525 (17)	0.0529 (16)	−0.0179 (16)	−0.0067 (15)	0.0033 (15)
C11	0.0533 (16)	0.0551 (17)	0.0497 (15)	−0.0173 (15)	0.0028 (14)	−0.0034 (14)

Geometric parameters (Å, º)

Cl1—C9	1.742 (3)	C3—H3A	0.9800
O1—C5	1.430 (3)	C4—C5	1.510 (3)
O1—C1	1.454 (3)	C4—H4A	0.9800
O2—C3	1.421 (3)	C5—H5A	0.9700
O2—H2C	0.8200	C5—H5B	0.9700
O3—C4	1.436 (3)	C6—C11	1.391 (4)
O3—H3B	0.8200	C6—C7	1.397 (4)
N1—C6	1.380 (3)	C7—C8	1.382 (4)
N1—C1	1.411 (3)	C7—H7A	0.9300
N1—H1B	0.9195	C8—C9	1.364 (4)

C1—C2	1.508 (3)	C8—H8A	0.9300
C1—H1A	0.9800	C9—C10	1.373 (4)
C2—C3	1.512 (3)	C10—C11	1.382 (4)
C2—H2A	0.9700	C10—H10A	0.9300
C2—H2B	0.9700	C11—H11A	0.9300
C3—C4	1.515 (3)		

C5—O1—C1	110.91 (18)	O3—C4—H4A	109.4
C3—O2—H2C	109.5	C5—C4—H4A	109.4
C4—O3—H3B	109.5	C3—C4—H4A	109.4
C6—N1—C1	124.9 (2)	O1—C5—C4	111.7 (2)
C6—N1—H1B	119.3	O1—C5—H5A	109.3
C1—N1—H1B	115.6	C4—C5—H5A	109.3
N1—C1—O1	109.19 (19)	O1—C5—H5B	109.3
N1—C1—C2	111.3 (2)	C4—C5—H5B	109.3
O1—C1—C2	109.23 (19)	H5A—C5—H5B	107.9
N1—C1—H1A	109.0	N1—C6—C11	119.7 (2)
O1—C1—H1A	109.0	N1—C6—C7	122.7 (2)
C2—C1—H1A	109.0	C11—C6—C7	117.6 (2)
C1—C2—C3	112.6 (2)	C8—C7—C6	121.0 (3)
C1—C2—H2A	109.1	C8—C7—H7A	119.5
C3—C2—H2A	109.1	C6—C7—H7A	119.5
C1—C2—H2B	109.1	C9—C8—C7	120.0 (3)
C3—C2—H2B	109.1	C9—C8—H8A	120.0
H2A—C2—H2B	107.8	C7—C8—H8A	120.0
O2—C3—C2	107.00 (19)	C8—C9—C10	120.5 (3)
O2—C3—C4	112.6 (2)	C8—C9—Cl1	120.2 (2)
C2—C3—C4	111.4 (2)	C10—C9—Cl1	119.3 (2)
O2—C3—H3A	108.6	C9—C10—C11	119.9 (3)
C2—C3—H3A	108.6	C9—C10—H10A	120.1
C4—C3—H3A	108.6	C11—C10—H10A	120.1
O3—C4—C5	108.14 (19)	C10—C11—C6	121.0 (3)
O3—C4—C3	111.5 (2)	C10—C11—H11A	119.5
C5—C4—C3	108.8 (2)	C6—C11—H11A	119.5

Hydrogen-bond geometry (Å, º)

D—H···*A*	*D*—H	H···*A*	*D*···*A*	*D*—H···*A*
O2—H2*C*···O3^{i}	0.82	1.93	2.739 (2)	170
O3—H3*B*···O1^{i}	0.82	1.98	2.797 (2)	175
N1—H1*B*···O2^{ii}	0.92	2.08	2.994 (3)	173

Symmetry codes: (i) −*x*+1, *y*+1/2, −*z*+1/2; (ii) −*x*, *y*−1/2, −*z*+1/2.