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Absolute Configuration of (+)-1-Phenyl-1,2,3,4-tetrahydroisoquinoline Hydrochloride

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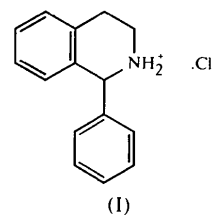
Abstract

(+)-1-Phenyl-1,2,3,4-tetrahydroisoquinoline with an $[\alpha]_D^{25}$ value of $+47.58^\circ$ in CCl_4 (c 2.83) has been synthesized and the crystal structure of its hydrochloride, $\text{C}_{15}\text{H}_{16}\text{N}^+\text{Cl}^-$, determined by X-ray methods. The

anomalous dispersion effect indicates that the absolute configuration is *S*.

Comment

1-Phenyl-1,2,3,4-tetrahydroisoquinoline (PTIQ) is a basic compound for the development of new drugs. The absolute configuration of (–)-PTIQ has been assigned to be *S* (Yamato *et al.*, 1990). However, our related compound suggested that (+)-PTIQ has the *S* configuration (Nakahara *et al.*, 1997). In order to confirm the absolute configuration, an X-ray diffraction study of the title compound, (I), was undertaken.



The optically resolved PTIQ we prepared had an $[\alpha]_D^{25}$ value of $+47.58^\circ$ in CCl_4 (c 2.83) indicating that it is the enantiomer of (–)-PTIQ (Leithe, 1929; Yamato *et al.*, 1990). Compound (I) crystallizes in the monoclinic space group $P2_1$ and the asymmetric unit contains two independent molecules, *A* and *A'*, having almost the same three-dimensional structure, with an r.m.s. deviation of 0.026 Å. They are related by a pseudo-twofold symmetry; their fractional atomic coordinates are related by the following equations: $x = -0.9995x' - 0.0117y' - 0.0389z' - 0.4600$, $y = -0.0027x' + 0.9999y' - 0.0330z' - 0.4185$, $z =$

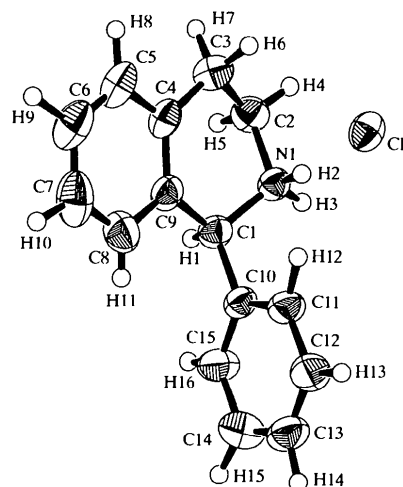


Fig. 1. ORTEPII drawing (Johnson, 1976) of molecule *A* of (I) with the atomic numbering. Displacement ellipsoids are shown at the 50% probability level and H atoms are shown as spheres of arbitrary size.

$0.0018x' - 0.0056y' - 1.0003z' + 1.5038$. The absolute configuration was unambiguously determined to be *S* by two methods, an *R*-factor ratio test (Hamilton, 1964) and a comparison between $|F_o(+)| - |F_o(-)|$ and $|F_c(+)| - |F_c(-)|$. The ratio between $wR(1S) = 0.0825$ and $wR(1R) = 0.1196$ indicates the *1S* configuration at more than the 99.9% significance level. The signs of the $|F_o(+)| - |F_o(-)|$ values for 40 selected Bijvoet mates were also consistent with those of $|F_c|$ values calculated for the *S* enantiomer. From these results, it was concluded that the absolute configuration of (-)-PTIQ should be *R*. Fig. 1 shows the molecular structure of molecule *A*, together with the absolute configuration. The bond lengths and angles given in Table 1 are within the normal ranges and the piperidine rings exhibit twist-chair conformations similar to those found for 1-methyl-1,2,3,4-tetrahydroisoquinoline (Ohkubo *et al.*, 1996). The interplanar angle between the least-squares planes of the benzene rings is 87.3(2)° for molecule *A* and 87.6(2)° for molecule *A'*.

The crystal packing is shown in Fig. 2. In both *A* and *A'*, the N1 atom links the two chloride ions related by a translation along the *a* axis through the two hydrogen bonds to form an infinite chain.

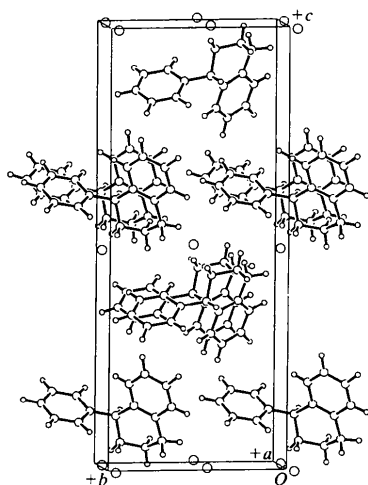


Fig. 2. Packing diagram of the crystal structure of (I).

Experimental

(+)-PTIQ was prepared and optically resolved according to the method of Leithe (1929), and its optical rotation was measured in CCl₄ solution (*c* 2.83) at 298 K using a 1 dm cell with a HORIBA polarimeter SEPA-200. Compound (I) was recrystallized from aqueous solution.

Crystal data

C₁₅H₁₆N⁺.Cl⁻
M_r = 245.751

Cu K α radiation
 $\lambda = 1.54184 \text{ \AA}$

Monoclinic
*P*2₁
a = 5.264 (3) \AA
b = 10.084 (4) \AA
c = 24.758 (3) \AA
 $\beta = 93.02 (4)^\circ$
V = 1312.4 (8) \AA^3
Z = 4
D_x = 1.244 Mg m⁻³
D_m not measured

Data collection

Rigaku AFC-7R diffractometer
 ω -2 θ scans
 Absorption correction:
 ψ scans (North *et al.*, 1968)
 $T_{\min} = 0.571$, $T_{\max} = 0.867$
 5824 measured reflections
 2636 independent reflections

Refinement

Refinement on *F*
R = 0.059
wR = 0.077
S = 1.160
 2627 reflections
 307 parameters
 Only coordinates of H atoms refined
 $w = 1/[\sigma^2(F_o) + 0.00286|F_o|^2]$
 $(\Delta/\sigma)_{\max} = 0.003$
 $\Delta\rho_{\max} = 0.28 \text{ e \AA}^{-3}$
 $\Delta\rho_{\min} = -0.34 \text{ e \AA}^{-3}$

Cell parameters from 20 reflections
 $\theta = 22.5\text{--}30.0^\circ$
 $\mu = 2.37 \text{ mm}^{-1}$
T = 298.2 K
 Plate
 0.62 × 0.28 × 0.06 mm
 Colorless

2627 reflections with *I* > 0
 $R_{\text{int}} = 0.047$
 $\theta_{\text{max}} = 70.0^\circ$
 $h = -6 \rightarrow 6$
 $k = -12 \rightarrow 12$
 $l = -30 \rightarrow 30$
 3 standard reflections every 150 reflections
 intensity decay: 3.95%

Extinction correction:
 Zachariasen (1967) type
 2 Gaussian isotropic
 Extinction coefficient:
 0.03 (2)
 Scattering factors from *International Tables for Crystallography* (Vol. C)
 Absolute structure:
 determined by *R*-factor ratio test (Hamilton, 1964) and the comparison between $|F_o(+)| - |F_o(-)|$ and $|F_c(+)| - |F_c(-)|$

Table 1. Selected geometric parameters (\AA , $^\circ$)

N1—C1	1.493 (4)	N1'—C1'	1.502 (5)
N1—C2	1.492 (5)	N1'—C2'	1.505 (6)
C1—C9	1.522 (5)	C1'—C9'	1.535 (6)
C1—C10	1.506 (5)	C1'—C10'	1.513 (6)
C2—C3	1.502 (6)	C2'—C3'	1.498 (6)
C3—C4	1.506 (6)	C3'—C4'	1.507 (6)
C4—C5	1.397 (6)	C4'—C5'	1.402 (6)
C4—C9	1.396 (6)	C4'—C9'	1.401 (6)
C5—C6	1.379 (8)	C5'—C6'	1.381 (8)
C6—C7	1.382 (9)	C6'—C7'	1.383 (10)
C7—C8	1.377 (8)	C7'—C8'	1.373 (8)
C8—C9	1.399 (5)	C8'—C9'	1.405 (6)
C10—C11	1.389 (5)	C10'—C11'	1.394 (6)
C10—C15	1.400 (5)	C10'—C15'	1.379 (6)
C11—C12	1.396 (6)	C11'—C12'	1.391 (7)
C12—C13	1.375 (7)	C12'—C13'	1.386 (8)
C13—C14	1.381 (8)	C13'—C14'	1.363 (9)
C14—C15	1.371 (7)	C14'—C15'	1.405 (9)
C1—N1—C2	111.6 (3)	C1'—N1'—C2'	110.8 (3)
N1—C1—C9	109.1 (3)	N1'—C1'—C9'	108.7 (3)
N1—C1—C10	111.0 (3)	N1'—C1'—C10'	110.0 (3)
C9—C1—C10	114.3 (3)	C9'—C1'—C10'	115.3 (3)
N1—C2—C3	110.0 (3)	N1'—C2'—C3'	110.6 (3)
C2—C3—C4	112.9 (3)	C2'—C3'—C4'	113.2 (4)
C3—C4—C5	119.0 (4)	C3'—C4'—C5'	119.2 (4)
C3—C4—C9	121.8 (4)	C3'—C4'—C9'	121.5 (4)

C5—C4—C9	119.1 (4)	C5'—C4'—C9'	119.2 (4)
C4—C5—C6	121.0 (5)	C4'—C5'—C6'	120.9 (5)
C5—C6—C7	120.1 (4)	C5'—C6'—C7'	119.7 (5)
C6—C7—C8	119.5 (5)	C6'—C7'—C8'	120.4 (5)
C7—C8—C9	121.3 (5)	C7'—C8'—C9'	121.0 (5)
C1—C9—C4	121.3 (3)	C1'—C9'—C4'	121.7 (4)
C1—C9—C8	119.7 (4)	C1'—C9'—C8'	119.5 (4)
C4—C9—C8	118.9 (4)	C4'—C9'—C8'	118.8 (4)
C1—C10—C11	121.8 (3)	C1'—C10'—C11'	121.3 (4)
C1—C10—C15	120.5 (3)	C1'—C10'—C15'	120.0 (4)
C11—C10—C15	117.7 (4)	C11'—C10'—C15'	118.7 (5)
C10—C11—C12	121.2 (4)	C10'—C11'—C12'	120.8 (4)
C11—C12—C13	119.4 (4)	C11'—C12'—C13'	119.5 (5)
C12—C13—C14	120.4 (4)	C12'—C13'—C14'	120.5 (5)
C13—C14—C15	120.0 (4)	C13'—C14'—C15'	119.8 (5)
C10—C15—C14	121.3 (4)	C10'—C15'—C14'	120.6 (5)

Table 2. Hydrogen-bonding geometry (Å, °)

D—H...A	D—H	H...A	D...A	D—H...A
N1—H2...C11	0.90	2.23	3.126 (3)	172
N1—H3...C11 ⁱ	1.01	2.09	3.084 (3)	167
N1'—H2'...C11'	0.90	2.22	3.116 (4)	176
N1'—H3'...C11' ⁱⁱ	1.11	1.98	3.074 (4)	168

Symmetry codes: (i) $1 + x, y, z$; (ii) $x - 1, y, z$.

The intensity data for hkl and $\bar{h}\bar{k}l$ were alternately measured every five reflections during the data collection. After structure determination, 40 reflections with large differences in F_c values between the Bijvoet reflections were chosen and their intensity data were measured more precisely. The structure was solved by direct methods and all non-H atoms were refined anisotropically by full-matrix least-squares techniques. Although all H atoms were located on a difference Fourier map, their positions were not refined. Their displacement parameters were assumed to be 1.2 times B_{eq} of the attached atom. For the R -factor ratio test, all atoms were refined again by full-matrix least-squares techniques with the reflections hkl and $\bar{h}\bar{k}l$ (total of 4919 reflections). The angles of the least-squares planes were calculated using the program LISTUP (Takenaka, 1990) and the other structural parameters were calculated using the program TEXSAN (Molecular Structure Corporation, 1997).

Data collection: Rigaku/AFC Diffractometer Control Software (Rigaku Corporation, 1995). Cell refinement: Rigaku/AFC Diffractometer Control Software. Data reduction: TEXSAN. Program(s) used to solve structure: SIR92 (Altomare *et al.*, 1994). Program(s) used to refine structure: TEXSAN. Molecular graphics: ORTEPII (Johnson, 1976). Software used to prepare material for publication: TEXSAN.

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The Monoclinic Form of Acetaminophen at 150 K

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Abstract

The crystal structure of the anti-inflammatory agent acetaminophen [*N*-(4-hydroxyphenyl)acetamide, C₈H₉NO₂] has been refined using a Siemens SMART three-circle automatic diffractometer with a charge-coupled device (CCD) detector at 150 K. Monoclinic crystals of acetaminophen were grown from ethanol solution (m.p. 443–444 K). The crystal structure exhibits two kinds of hydrogen bonds [2.656(2) Å, OH donates to O=C; 2.914(2) Å, HO accepts from HN]. The molecules form a pleated sheet parallel to the (101) plane. The sheets are stacked along [010]. The molecules within a stack are held together by van der Waals interactions. Molecules from different sheets form 'head-to-tail'-type dimers.

Comment

Acetaminophen, (I), is the most prominent pain-relieving drug among acetanilide derivatives. Its physical and chemical properties have been intensively studied by Grant & Chow (1991). Precise data on its crystal structure are required for controlling processes such as dissolution and sublimation, which are important