C13—C14	1.398 (13)	1.384 (13)
C14—C15	1.391 (13)	1.393 (13)
C15—C16	1.408 (13)	1.404 (13)
C16—C17	1.362 (14)	1.372 (13)
C1—C2 C2—C3 C2—O1	Molecule <i>C</i> 1.50 (2) 1.50 (2) 1.44 (1)	Molecule <i>D</i> 1.50(2)*/1.49(3)† 1.49(2)/1.47(4) 1.51(2)/1.38(4)

* Most occupied position. † Least occupied position.

Table 3. Hydrogen-bonding geometry (Å, °)

$D - H \cdots A$	D—H	$\mathbf{H} \cdot \cdot \cdot \mathbf{A}$	$D \cdots A$	$D = H \cdot \cdot \cdot A$
N1A—H1A···O3A ⁱ	0.91	2.11	2.89(1)	143
NIA—HIA···O4A ⁱ	0.91	2.30	3.14(1)	152
N1AH2A···O3B	0.91	1.86	2.77 (1)	175
$N1A - H3A \cdot \cdot \cdot O1D^{ii}$	0.91	1.97	2.81(1)	152
$N2A - H6A \cdot \cdot \cdot O2B^{iii}$	0.88	2.09	2.95(1)	165
N3A—H17A···O1B	0.88	2.12	2.97 (1)	163
$N1B - H1B \cdot \cdot \cdot O3B^{iv}$	0.91	2.32	3.03(1)	134
$N1B$ — $H1B \cdot \cdot \cdot O4B^{iv}$	0.91	1.91	2.74 (1)	151
N1 <i>B</i> —H2 <i>B</i> ···O3A	0.91	2.11	2.80(1)	132
N1 <i>B</i> —H3 <i>B</i> ···O1 <i>C</i>	0.91	2.05	2.80(1)	140
$N2B - H6B \cdot \cdot \cdot O2A^{v}$	0.88	2.00	2.84 (1)	159
N3BH17B···O1A	0.88	2.18	3.05(1)	172
$O5A - H25A \cdot \cdot \cdot O1E^{vi}$	0.84	2.01	2.84(1)	171
O5B—H25B· · · O1G	0.84	2.02	2.70(1)	137
O1 <i>C</i> —H8 <i>C</i> ···O1 <i>H</i>	0.84	2.04	2.83 (1)	157
01 <i>E</i> —H1 <i>E</i> ···O1 <i>F</i>	1.05	2.11 (12)	2.89(1)	128 (9)
01 <i>G</i> —H1 <i>G</i> ···O2A	0.97	2.34 (17)	2.96(1)	121 (13)
Symmetry codes: (i) $x, y, z - 1$; (ii) $x - 1, y, z - 1$; (iii) $x - 1, y, z$;				

(iv) x, y, 1 + z; (v) 1 + x, y, z; (vi) x, y - 1, z.

Table 4. Molecular conformations (°) for Gly-L-Leu-L-Tyr in crystal structures

Torsion	2-Propan	ol solvate*	DMS	O solvate†	Cu co	omplex‡
angle	Α	В	Α	В	Α	В
Backbone	es§					
ψ_1	156.6(7)	169.5 (7)	154	170	-176	180
ω_1	177.7 (7)	178.5 (7)	-179	177	-177	-173
φ_2	-149.3 (7)	-123.5 (8)	-146	-132	-131	-130
ψ_2	132.7 (7)	143.6(7)	135	148	142	139
ω_2	177.5 (7)	173.1 (7)	176	174	177	-177
φ_3	-129.7 (8)	-145.5(7)	-140	-146	-142	- 149
ψ_T	-13.4 (10)	-6.7(11)	-10	-10	-23	-20
Side chai	ns					
χ_{7}^{1}	179.5 (7)	-55.0(10)	-176	-60	176	-53
$\chi_{2}^{2,1}$	153.1 (9)	169.1 (8)	151	169	153	174
$x_2^{2,2}$	-83.8(10)	-67.6(10)	86	-71	-92	-67
χ_3^{-1}	65.0 (9)	65.2 (9)	59	66	58	54
$\chi_{3}^{2,1}$	86.2 (9)	-86.8 (9)	88	-88	84	-79
$\chi_{3}^{2,2}$	-94.8(9)	98.0 (9)	- 89	99	-86	90

* This work. † Subramanian & Parthasarathy (1987). ‡ Franks & van der Helm (1970). § $\psi_1 = N1-C1-C2-N2$, $\omega_1 = C3-N2-C2-C1$, $\varphi_2 = C2-N2-C3-C8$, $\psi_2 = N2-C3-C8-N3$, $\omega_2 = C9-N3-C8-C3$, $\varphi_3 = C8-N3-C9-C10$, $\psi_T = N3-C9-C10-O3$, $\chi_2^{-1} = N2-C3-C4-C5$, $\chi_2^{2,1} = C3-C4-C5-C6$, $\chi_2^{-2,2} = C3-C4-C5-C7$, $\chi_3^{-1} = N3-C9-C11-C12$, $\chi_3^{-2,1} = C9-C11-C12-C13$, $\chi_3^{-2,2} = C9-C11-C12-C17$.

The structure was solved by direct methods using SIR92 (Altomare *et al.*, 1992) and refined with SHELXL93 (Sheldrick, 1993). All heavy atoms, except C and amide N backbone atoms and the disordered 2-propanol C atoms, were refined anisotropically. H atoms with known geometry were added in theoretical positions. The coordinates for the H atoms

© 1996 International Union of Crystallography Printed in Great Britain – all rights reserved 'ride' on the coordinates of the corresponding heavy atom. Isotropic temperature factors for amino, hydroxyl and methyl H atoms were fixed to 1.5 times U_{eq} of the bonded heavy atom. Isotropic temperature factors for the remaining H atoms were fixed to 1.2 times U_{eq} of the attached heavy atom. The coordinates of the hydroxyl H atoms for the two Tyr residues and the two 2-propanol molecules were refined using a 'rotating group refinement' constraint with O—H = 0.84 Å. Three of the H atoms in the water molecules were located from the difference Fourier map, and their positions refined. Two alternative positions have been refined for the C atoms C1D and C2D, with occupancy factors 0.66 (3) for the pair C1D2 and C2D2, and 0.34 (3) for C1D1 and C2D1.

Lists of structure factors, anisotropic displacement parameters, Hatom coordinates and complete geometry have been deposited with the IUCr (Reference: AB1338). Copies may be obtained through The Managing Editor, International Union of Crystallography, 5 Abbey Square, Chester CH1 2HU, England.

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1-Phenyl-2-(propylamino)pentane, a Memory Enhancer

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Abstract

The absolute configuration of [(R)-1-phenyl-2-pentyl]propylammonium chloride, $C_{14}H_{24}N^+$.Cl⁻, has been determined. The compound is a memory enhancer. Since the pharmacological effect of the related compound selegiline is highly dependent upon its configuration, the title compound may be expected to behave in a similar manner.

Comment

Selegiline has been widely used in the treatment of Parkinson's disease. A number of similar compounds have been synthesized over the past few years. Of these we now describe the structure of [(R)-1-phenyl-2pentyllpropylammonium chloride, (1), which is reported to facilitate learning and retention (Knoll, Knoll, Török, Timár & Yasar, 1992; Knoll & Miklya, 1994). Whereas amphetamines facilitate performance in a very narrow low dosage range which changes at a modest elevation of the dose into the opposite effect, compound (1) improves performance within a reasonably broad range. In the case of selegiline, the R enantiomer has much better pharmacological properties than its S counterpart (Robinson, 1985; Magyar, Vizi, Ecsery & Knoll, 1967). Therefore, we believe that an accurate characterization of these compounds is vitally important.



The crystal structure of (1) contains two crystallographically independent molecules in the asymmetric unit (Fig. 1). Both of these molecules are in the R configuration. The conformations of the two species are different, however, particularly with regard to the aliphatic side chains (Fig. 2). As far as the crystal packing is concerned, it is worth noting that parallel chains of hydrogen bonds formed by the chloride anions and the alkylammonium cations pack into hydrophilic layers in the *ab* plane (Fig. 3). This layer-like hydrogen-bond network is -probably the most important contributor to intermolecular interactions. In between the hydrophilic layers there are hydrophobic layers packed with phenyl groups in the middle (Fig. 3).



Fig. 1. The molecular structure and atomic numbering for (1). Displacement ellipsoids are plotted at the 50% probability level.



Fig. 2. Superposition of the two conformationally different molecules in the asymmetric unit.



Fig. 3. A packing diagram for compound (1).

Experimental

The title compound was prepared according to Knoll, Knoll, Török, Timár & Yasar (1992) and Knoll *et al.* (1988, 1989).

Crystal data

C₁₄H₂₄N⁺.Cl⁻ $M_r = 241.79$ Monoclinic $P2_1$ a = 7.665 (1) Å b = 14.816 (1) Å c = 13.237 (1) Å $\beta = 93.830 (1)^\circ$ $V = 1500.0 (2) Å^3$ Z = 4 $D_x = 1.071 \text{ Mg m}^{-3}$ D_m not measured Cu K α radiation $\lambda = 1.5418$ Å Cell parameters from 25 reflections $\theta = 41.05-49.80^{\circ}$ $\mu = 2.050 \text{ mm}^{-1}$ T = 296 (2) K Plate $0.25 \times 0.20 \times 0.11 \text{ mm}$ Transparent

Data collection	
Enraf–Nonius CAD-4	$\theta_{\rm max} = 75.04^{\circ}$
diffractometer	$h = 0 \rightarrow 9$
$\omega/2\theta$ scans	$k = -18 \rightarrow 0$
Absorption correction:	$l = -16 \rightarrow 16$
none	3 standard reflections
3222 measured reflections	monitored every 150
3222 independent reflections	reflections
2186 observed reflections	intensity decay: 1.2%
$[I>2\sigma(I)]$	
Refinement	
Refinement on F^2	Extinction correction:
R(F) = 0.0528	SHELXL93 (Sheldrick,
$wR(F^2) = 0.1654$	1993)
S = 1.024	Extinction coefficient:
3219 reflections	0.0029 (10)
295 parameters	Atomic scattering factors
Only H-atom U's refined	from International Tables
$w = 1/[\sigma^2(F_{\rho}^2) + (0.1161P)^2]$	for Crystallography (1992,
where $P = (F_o^2 + 2F_c^2)/3$	Vol. C, Tables 4.2.6.8 and
$(\Delta/\sigma)_{\rm max} = 0.013$	6.1.1.4)
$\Delta \rho_{\rm max} = 0.482 \ {\rm e} \ {\rm \AA}^{-3}$	Absolute configuration:
$\Delta \rho_{\rm min} = -0.264 \ {\rm e} \ {\rm \AA}^{-3}$	Flack (1983) parameter

Table 1. Fractional atomic coordinates and equivalent isotropic displacement parameters (Å²)

 $U_{\rm eq} = (1/3) \sum_i \sum_j U_{ij} a_i^* a_i^* \mathbf{a}_i \cdot \mathbf{a}_j.$

= 0.00(3)

	x	у	z	U_{eq}
C11	-0.56763 (12)	1.12060 (7)	1.02645 (9)	0.0655 (3)
NI	-0.9531 (4)	1.1664 (3)	1.0732(3)	0.0575 (8)
C1	-1.1422 (10)	1.1485 (6)	0.8000 (5)	0.106 (2)
C2	-1.0233(8)	1.1790 (5)	0.8896 (4)	0.088 (2)
C3	-1.0657 (6)	1.1300 (4)	0.9856 (3)	0.0668 (11)
C4	-0.9866(6)	1.1320 (3)	1.1754 (3)	0.0620(11)
C5	-1.1690 (6)	1.1636(4)	1.2017 (3)	0.0736 (13)
C6	-1.2084(6)	1.1480 (3)	1.3109 (4)	0.0638 (11)
C7	-1.3463 (7)	1.0945 (4)	1.3353 (5)	0.082 (2)
C8	-1.3804 (10)	1.0821 (5)	1.4349 (6)	0.104 (2)
C9	-1.2815 (10)	1.1250(7)	1.5107 (5)	0.102 (2)
C10	-1.1491 (9)	1.1785 (6)	1.4873 (4)	0.096 (2)
C11	-1.1100(7)	1.1898 (5)	1.3882 (4)	0.0782 (14)
C12	-0.9539(9)	1.0311 (4)	1.1845 (5)	0.094 (2)
C13	-0.8961 (15)	0.9990 (9)	1.2907 (8)	0.163 (5)
C14	-0.8464(17)	0.9069 (8)	1.3015 (10)	0.175 (5)
C11'	-0.96329 (15)	0.87549 (8)	0.92696 (10)	0.0736 (3)
N1'	-0.5602 (5)	0.9271 (2)	0.9280(3)	0.0580 (8)
C1′	-0.4018 (13)	0.8230(6)	1.1814 (5)	0.117 (3)
C2′	-0.5212 (9)	0.8765 (6)	1.1046 (4)	0.093 (2)
C3′	-0.4514 (7)	0.8723 (4)	1.0033 (4)	0.0763 (13)
C4′	-0.4760 (7)	0.9362 (3)	0.8285 (4)	0.0638 (11)
C5′	-0.4508 (8)	0.8451 (4)	0.7806 (4)	0.0748 (13)
C6′	-0.3715 (7)	0.8500(3)	0.6785 (4)	0.0719(13)
C7′	-0.2116 (7)	0.8866 (4)	0.6684 (4)	0.0813 (14)
C8′	-0.1411 (9)	0.8926 (5)	0.5754 (5)	0.096 (2)
C9′	-0.2339 (12)	0.8588 (6)	0.4915 (6)	0.108 (2)
C10′	-0.3914 (12)	0.8215 (6)	0.5004 (5)	0.107 (2)
C11'	-0.4602 (10)	0.8179 (5)	0.5940 (4)	0.093 (2)
C12′	-0.5500 (13)	1.0115 (7)	0.7657 (6)	0.121 (3)
C13′	-0.7091 (12)	0.9961 (9)	0.7268 (9)	0.146 (4)
C14′	-0.7691 (12)	1.0853 (8)	0.6457 (7)	0.137 (3)

Table 2. Selected geometric parameters (Å, °)

N1C4	1.484 (6)	N1'C3'	1.496 (6)
N1C3	1.499 (6)		1.511 (6)
C4-N1-C3	117.2 (4)	C3'—N1'—C4'	112.2 (4)

C1-C2-C3-N1	-176.5 (5)
C4-N1-C3-C2	174.9 (4)
C3N1C4C5	-65.8 (5)
C12-C4-C5-C6	64.0 (7)
N1-C4-C5-C6	-169.6 (5)
C5-C4-C12-C13	-84.4 (8)
C4C12C13C14	-173.7 (9)
C1'-C2'-C3'-N1'	-177.5 (6)
C4'-N1'-C3'-C2'	170.8 (5)
C3'-N1'-C4'-C5'	60.7 (6)
C12'-C4'-C5'-C6'	44.2 (9)
N1'-C4'-C5'-C6'	178.3 (4)
C5'-C4'-C12'-C13'	61.9 (10)
C4'-C12'-C13'-C14'	-174.4 (6)

Data collection: CAD-4 Software (Enraf-Nonius, 1989). Cell refinement: CAD-4 Software. Data reduction: CAD-4 Software. Program(s) used to solve structure: SHELXS86 (Sheldrick, 1990). Program(s) used to refine structure: SHELXL93 (Sheldrick, 1993). Molecular graphics: TEXSAN (Molecular Structure Corporation, 1985, 1992). Software used to prepare material for publication: SHELXL93.

Lists of structure factors, anisotropic displacement parameters, Hatom coordinates, complete geometry and torsion angles have been deposited with the IUCr (Reference: KA1192). Copies may be obtained through The Managing Editor, International Union of Crystallography, 5 Abbey Square, Chester CH1 2HU, England.

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