

The absolute configuration of a new amino-alkylphenol derived from (*R*)-(-)-2-phenylglycine

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## Key indicators

Single-crystal X-ray study  
*T* = 298 K  
Mean  $\sigma(\text{C}-\text{C}) = 0.005 \text{ \AA}$   
*R* factor = 0.052  
*wR* factor = 0.144  
Data-to-parameter ratio = 14.7For details of how these key indicators were automatically derived from the article, see <http://journals.iucr.org/e>.

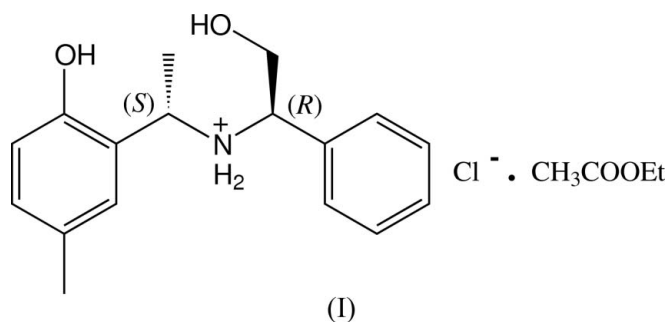
The chiral aminoalkylphenol *N*-[(*S*)-1-(5-methyl-2-hydroxyphenyl)ethyl]-*N*-[(*R*)-2-hydroxy-1-phenylethyl]amine,  $\text{C}_{17}\text{H}_{21}\text{NO}_2$ , was synthesized starting from *R*-(-)-2-phenylglycine. The corresponding hydrochloride salt was prepared and crystallized as an ethyl acetate solvate, affording *N*-[(*R*)-2-hydroxy-1-phenylethyl]-*N*-[(*S*)-1-(5-methyl-2-hydroxyphenyl)ethyl]-aminium chloride ethyl acetate solvate,  $\text{C}_{17}\text{H}_{22}\text{NO}_2^+\cdot\text{Cl}^-\cdot\text{C}_4\text{H}_8\text{O}_2$ , and the absolute configuration of the new stereogenic centre determined, from which that of the chiral aminoalkylphenol was inferred. The dihedral angle between the two aromatic rings is  $40.74 (11)^\circ$ .

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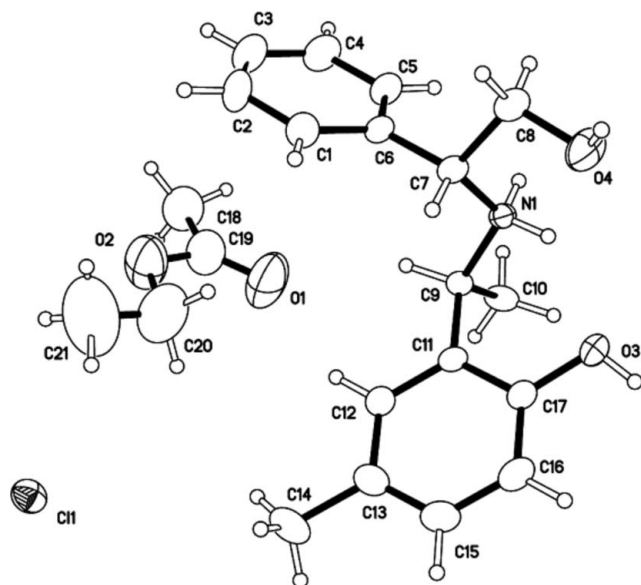
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## Comment

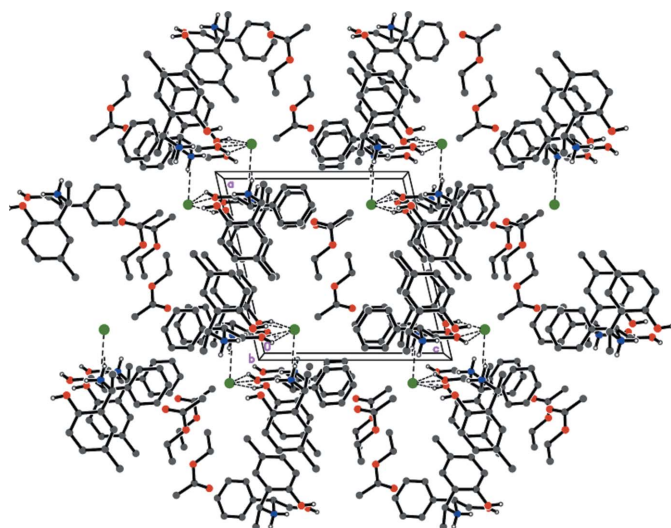
The synthesis of new chiral ligands is widespread in asymmetric synthesis (Cimarelli & Palmieri, 1998, 2000; Tseng & Yang, 2004; Tümerdem *et al.*, 2005). Among them, enantiopure aminoalkylphenols have attracted wide attention for the reason that they can be used in catalytic asymmetric reactions (Puigjaner *et al.*, 1999; Guangyou *et al.*, 2003; Li *et al.*, 2004; Watts *et al.*, 2005), which is one of the most active areas of research in organic chemistry (Joshi & Malhotra, 2003). However, the chiral ligand is not always readily or economically available. Most of these aminoalkylphenols are derived from readily available natural products (Soai & Niwa, 1992). Amino acids can be converted into the corresponding aminoalcohols by reduction with  $\text{NaBH}_4$  in THF. As the reduction method gave a single enantiomer in high yield (McKennon *et al.*, 1993), in our preliminary work we chose *R*-(-)-2-phenylglycine as starting material.



The aminoalkylphenol was prepared by conventional condensation of 2-hydroxy-5-methylacetophenone with *R*-(-)-phenylglycinol, followed by reduction with sodium borohydride (see *Experimental*). The *R,S* diastereoisomer is the main product, in agreement with the fact that attack of the hydride occurs on the less-hindered face. In order to confirm the structure of this aminoalkylphenol, the corresponding



**Figure 1**  
The asymmetric unit of (I), with displacement ellipsoids drawn at the 30% probability level for non-H atoms.



**Figure 2**  
A packing diagram for (I), viewed down the *b* axis. Hydrogen bonds are shown as dashed lines. H atoms not involved in hydrogen bonding have been omitted.

hydrochloride ethyl acetate solvate, (I), was prepared and an X-ray crystallographic study carried out (Fig. 1). Geometric parameters (Table 1) are similar to those found in a closely related aminoalkylphenol (Yang *et al.*, 2005; Zhang *et al.*, 2006).

The absolute configuration of (I) is found to be *R,S*, from which that of its neutral aminoalkylphenol precursor is inferred to be the same. The dihedral angle between the planes of the C1–C6 and C11–C17 aromatic rings is 40.74 (11)°. Ethyl acetate is included as a solvent molecule. Intermolecular O–H···Cl and N–H···Cl (Fig. 2), as well as intramolecular N–H···O hydrogen bonds are formed, thereby stabilizing the crystal structure (Table 2).

## Experimental

*R*-(–)-2-Phenylglycinol was prepared by the reduction of *R*-(–)-2-phenylglycine with NaBH<sub>4</sub> in THF [80.2% yield,  $[\alpha]_D^{24} = -25.5$  (*c* 6, MeOH)] (Abiko & Masamune, 1992; McKennon *et al.*, 1993; Demir *et al.*, 1999). *R*-(–)-2-Phenylglycinol (2 mmol) and 1-(2-hydroxy-4-methylphenyl)-ethanone (2 mmol) were dissolved in methanol (15 ml) and reacted at 298 K for 12 h. The solvent was then removed by evaporation and THF/ethanol (15 ml, 1:1 *v/v*) and NaBH<sub>4</sub> (8 mmol) were added at 273 K. The reaction was quenched with 5 M HCl and then neutralized with NaOH. The organic layer was dried with anhydrous Na<sub>2</sub>SO<sub>4</sub>, filtered, and the solvent was removed under reduced pressure. Further purification was carried out by thin-layer silica-gel chromatography [chloroform/methanol 40:1; 70.1% yield;  $[\alpha]_D^{27} = -77.4$  (*c* 1/2, CHCl<sub>3</sub>)]. The hydrochloride was finally obtained by reaction of the amine (0.1 mmol) dissolved in methanol (10 ml) with concentrated HCl (0.1 ml) at 298 K, affording a white solid. Compound (I) was crystallized from a 2-propanol/ethyl acetate (1:20 *v/v* mixture (83% yield).

## Crystal data

C<sub>17</sub>H<sub>22</sub>NO<sub>2</sub><sup>+</sup>·Cl<sup>−</sup>·C<sub>4</sub>H<sub>8</sub>O<sub>2</sub>  
*M<sub>r</sub>* = 395.91  
 Monoclinic, *P*2<sub>1</sub>  
*a* = 12.043 (2) Å  
*b* = 7.7441 (16) Å  
*c* = 12.043 (2) Å  
 $\beta$  = 103.07 (2)°  
*V* = 1094.2 (4) Å<sup>3</sup>

*Z* = 2  
*D<sub>x</sub>* = 1.202 Mg m<sup>−3</sup>  
 Mo *K*α radiation  
 $\mu$  = 0.20 mm<sup>−1</sup>  
*T* = 298 (2) K  
 Plate, colourless  
 0.50 × 0.40 × 0.12 mm

## Data collection

Bruker SMART diffractometer  
 $\varphi$  and  $\omega$  scans  
 Absorption correction: multi-scan  
 (SADABS; Sheldrick, 1996)  
*T<sub>min</sub>* = 0.907, *T<sub>max</sub>* = 0.977

5629 measured reflections  
 3672 independent reflections  
 3546 reflections with *I* > 2σ(*I*)  
*R<sub>int</sub>* = 0.025  
 $\theta_{\max}$  = 25.2°

## Refinement

Refinement on *F*<sup>2</sup>  
*R*[*F*<sup>2</sup> > 2σ(*F*<sup>2</sup>)] = 0.052  
*wR*(*F*<sup>2</sup>) = 0.144  
*S* = 1.05  
 3672 reflections  
 250 parameters  
 H-atom parameters constrained

$w = 1/[\sigma^2(F_o^2) + (0.0958P)^2 + 0.1694P]$   
 where  $P = (F_o^2 + 2F_c^2)/3$   
 $(\Delta/\sigma)_{\max} = 0.001$   
 $\Delta\rho_{\max} = 0.23 \text{ e \AA}^{-3}$   
 $\Delta\rho_{\min} = -0.23 \text{ e \AA}^{-3}$   
 Absolute structure: Flack (1983),  
 1550 Friedel pairs  
 Flack parameter: 0.00 (8)

**Table 1**

Selected geometric parameters (Å, °).

C6–C7	1.503 (4)	C9–N1	1.510 (3)
C7–N1	1.499 (4)	C9–C10	1.527 (4)
C7–C8	1.521 (4)	C13–C14	1.521 (5)
C9–C11	1.508 (4)	C17–O3	1.370 (4)
N1–C7–C6	112.4 (2)	N1–C9–C10	108.8 (2)
N1–C7–C8	107.5 (2)	O3–C17–C16	123.2 (3)
C6–C7–C8	112.2 (2)	O3–C17–C11	117.0 (2)
O4–C8–C7	107.9 (3)		

**Table 2**  
Hydrogen-bond geometry (Å, °).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
N1—H1A $\cdots$ O3	0.90	2.00	2.719 (3)	136
N1—H1A $\cdots$ O3	0.90	2.00	2.719 (3)	136
N1—H1B $\cdots$ Cl1 <sup>i</sup>	0.90	2.31	3.181 (2)	163
O3—H3 $\cdots$ Cl1 <sup>ii</sup>	0.82	2.21	3.022 (2)	172
O4—H4 $\cdots$ Cl1 <sup>iii</sup>	0.82	2.35	3.149 (3)	164

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

The cell obtained for (I) has  $a = c$ , although actually belonging to Laue class  $2/m$ . Such a case had been observed in previous reports (e.g. Sani *et al.*, 2005). In order to converge to a sensible geometry for the solvent molecule, the C20—C21 bond length was restrained to 1.50 (1) Å and atoms within this group were subjected to a rigid-bond restraint: the  $U^{ij}$  parameters in the direction of the bonds were restrained to be equal within an s.u. of 0.01 Å<sup>2</sup>. Finally, for the complete asymmetric unit, atoms closer than 1.7 Å were restrained to have the same  $U^{ij}$  components, within an s.u. of 0.02 Å<sup>2</sup>. All H atoms were placed in idealized positions and constrained to ride on their parent atom; constrained distances: O—H = 0.82 Å, N—H = 0.90 Å, and C—H = 0.93, 0.96, 0.97 and 0.98 Å for aromatic, methyl, methylene and methine groups, respectively. Isotropic displacement parameters were fixed at  $U_{\text{iso}}(\text{H}) = 1.5U_{\text{iso}}(\text{carrier atom})$  for OH and methyl groups and at  $U_{\text{iso}}(\text{H}) = 1.2U_{\text{iso}}(\text{carrier atom})$  for other groups.

Data collection: *SMART* (Bruker, 1997); cell refinement: *SAINTE* (Bruker, 1997); data reduction: *SAINTE*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *SHELXTL* (Bruker, 2001); software used to prepare material for publication: *SHELXTL*.

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