

Acta Crystallographica Section E

## Structure Reports

Online

ISSN 1600-5368

# N-[2-(2-Chlorophenyl)-2-hydroxyethyl]-propan-2-aminium chloride

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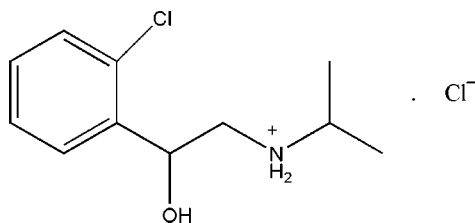
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Received 1 August 2009; accepted 6 August 2009

Key indicators: single-crystal X-ray study;  $T = 296$  K; mean  $\sigma(\text{C}-\text{C}) = 0.005$  Å;  $R$  factor = 0.032;  $wR$  factor = 0.107; data-to-parameter ratio = 21.3.

In the title compound,  $\text{C}_{11}\text{H}_{17}\text{ClNO}^+\cdot\text{Cl}^-$ , the side chain of the ethylamine group is orientated approximately perpendicular to the benzene ring, the dihedral angle between the C/C/N plane of the ethylamine group and the benzene plane being  $83.5(3)^\circ$ . In the crystal structure, intermolecular  $\text{O}-\text{H}\cdots\text{Cl}$  and  $\text{N}-\text{H}\cdots\text{Cl}$  hydrogen bonds are observed. The crystal studied was an inversion twin with a 0.51(10):0.49(10) domain ratio.

## Related literature

 For a related structure, see: Tang *et al.* (2009).


## Experimental

### Crystal data

 $\text{C}_{11}\text{H}_{17}\text{ClNO}^+\cdot\text{Cl}^-$ 
 $M_r = 250.16$ 

Orthorhombic,  $P2_12_12_1$   
 $a = 7.3460(3)$  Å  
 $b = 11.7721(5)$  Å  
 $c = 15.2377(8)$  Å  
 $V = 1317.72(10)$  Å<sup>3</sup>

$Z = 4$   
 Mo  $K\alpha$  radiation  
 $\mu = 0.47$  mm<sup>-1</sup>  
 $T = 296$  K  
 $0.40 \times 0.36 \times 0.32$  mm

### Data collection

Rigaku R-AXIS RAPID diffractometer  
 Absorption correction: multi-scan (ABSCOR; Higashi, 1995)  
 $T_{\min} = 0.835$ ,  $T_{\max} = 0.864$

12577 measured reflections  
 2977 independent reflections  
 1874 reflections with  $I > 2\sigma(I)$   
 $R_{\text{int}} = 0.031$

### Refinement

$R[F^2 > 2\sigma(F^2)] = 0.032$   
 $wR(F^2) = 0.107$   
 $S = 1.00$   
 2977 reflections  
 140 parameters  
 H-atom parameters constrained

$\Delta\rho_{\max} = 0.29$  e Å<sup>-3</sup>  
 $\Delta\rho_{\min} = -0.29$  e Å<sup>-3</sup>  
 Absolute structure: Flack (1983),  
 1243 Friedel pairs  
 Flack parameter: 0.51(10)

Table 1

Hydrogen-bond geometry (Å, °).

$D-\text{H}\cdots A$	$D-\text{H}$	$\text{H}\cdots A$	$D\cdots A$	$D-\text{H}\cdots A$
$\text{N1}-\text{H112}\cdots\text{Cl2}$	0.90	2.36	3.199(2)	156
$\text{O1}-\text{H1}\cdots\text{Cl2}^i$	0.82	2.33	3.143(2)	169
$\text{N1}-\text{H111}\cdots\text{Cl2}^{ii}$	0.90	2.28	3.138(2)	160

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

Data collection: *PROCESS-AUTO* (Rigaku, 2006); cell refinement: *PROCESS-AUTO*; data reduction: *CrystalStructure* (Rigaku/MS, 2007); program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *ORTEP-3 for Windows* (Farrugia, 1997); software used to prepare material for publication: *WinGX* (Farrugia, 1999).

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: IS2447).

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## supporting information

*Acta Cryst.* (2009). E65, o2187 [doi:10.1107/S1600536809031146]

***N*-[2-(2-Chlorophenyl)-2-hydroxyethyl]propan-2-aminium chloride****Bi-Wei Song, Lin-Jun Xie, Ling-Ling Dong, Zhan Tang and Hai Feng****S1. Comment**

The title compound (clorprenaline hydrochloride) is one of a series of structurally related  $\beta$ -adrenoceptorblocking drugs.

In the molecular structure (Fig. 1), there are no unusual bond distances or angles. The Cl atom and the phenyl plane is almost planar with the deviation of 0.0037 Å. The dihedral angle between the plane formed by C7/C8/N1 and the phenyl plane is 83.5 (3)°, which shows that the two planes are almost perpendicular. The C9—N1 distance of 1.506 Å is longer than the value of the similar bond distance of 1.474 Å (Tang *et al.*, 2009).

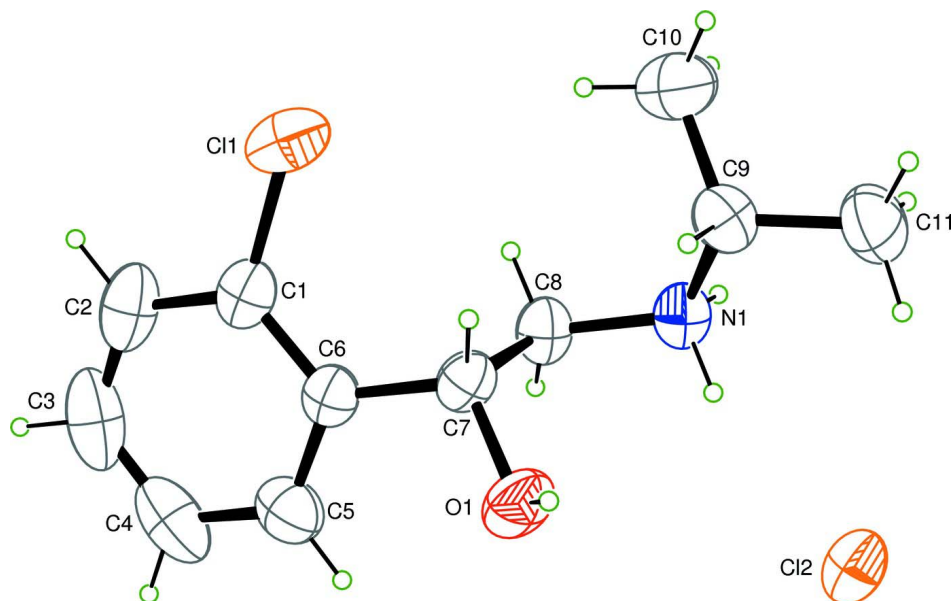
O—H...Cl and N—H...Cl hydrogen bonds are found in the crystal structure and are essential forces in crystal formation. The hydroxyl hydrogen at O1 acts as a donor to Cl2. The ethylamine hydrogens at N1 also act as donors to Cl2.

**S2. Experimental**

Racemic Clorprenaline hydrochloride was purchased from ShangHai Shengxin Medicine & Chemical Co., Ltd. ShangHai, China. Racemic Clorprenaline hydrochloride (5 g) was dissolved in ethanol (75 ml) and then hydrochloric acid was added to give pH of about 4. Colorless crystal of (I) separated from the solution in about 80% yield after one day.

**S3. Refinement**

All of the H atoms were placed in calculated positions and allowed to ride on their parent atoms, with C—H = 0.93 (aromatic), 0.98 (methine), 0.97 (methylene), 0.96 Å (methyl), O—H = 0.82 Å and N—H = 0.90 Å, and with  $U_{\text{iso}}(\text{H}) = 1.2\text{--}1.5$  times  $U_{\text{eq}}$  of the parent atoms.

**Figure 1**

The molecular structure of (I) with atom labels, showing 40% probability displacement ellipsoids.

### ***N*-[2-(2-Chlorophenyl)-2-hydroxyethyl]propan-2-aminium chloride**

#### *Crystal data*

$C_{11}H_{17}ClNO^+Cl^-$

$M_r = 250.16$

Orthorhombic,  $P2_12_12_1$

Hall symbol: P 2ac 2ab

$a = 7.3460$  (3) Å

$b = 11.7721$  (5) Å

$c = 15.2377$  (8) Å

$V = 1317.72$  (10) Å<sup>3</sup>

$Z = 4$

$F(000) = 528$

$D_x = 1.261$  Mg m<sup>-3</sup>

Mo  $K\alpha$  radiation,  $\lambda = 0.71073$  Å

Cell parameters from 8627 reflections

$\theta = 3.1$ – $27.4^\circ$

$\mu = 0.47$  mm<sup>-1</sup>

$T = 296$  K

Chunk, colorless

$0.40 \times 0.36 \times 0.32$  mm

#### *Data collection*

Rigaku R-AXIS RAPID  
diffractometer

Radiation source: rotating anode

Graphite monochromator

Detector resolution: 10.00 pixels mm<sup>-1</sup>

$\omega$  scans

Absorption correction: multi-scan

(*ABSCOR*; Higashi, 1995)

$T_{\min} = 0.835$ ,  $T_{\max} = 0.864$

12577 measured reflections

2977 independent reflections

1874 reflections with  $I > 2\sigma(I)$

$R_{\text{int}} = 0.031$

$\theta_{\max} = 27.4^\circ$ ,  $\theta_{\min} = 3.1^\circ$

$h = -9 \rightarrow 9$

$k = -15 \rightarrow 14$

$l = -19 \rightarrow 19$

#### *Refinement*

Refinement on  $F^2$

Least-squares matrix: full

$R[F^2 > 2\sigma(F^2)] = 0.032$

$wR(F^2) = 0.107$

$S = 1.00$

2977 reflections

140 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  
 $w = 1/[\sigma^2(F_o^2) + (0.0431P)^2 + 0.5P]$   
 where  $P = (F_o^2 + 2F_c^2)/3$   
 $(\Delta/\sigma)_{\max} < 0.001$   
 $\Delta\rho_{\max} = 0.29 \text{ e } \text{\AA}^{-3}$   
 $\Delta\rho_{\min} = -0.29 \text{ e } \text{\AA}^{-3}$

Extinction correction: *SHELXL97* (Sheldrick, 2008),  $F_c^* = kFc[1 + 0.001xFc^2\lambda^3/\sin(2\theta)]^{-1/4}$   
 Extinction coefficient: 0.0054 (12)  
 Absolute structure: Flack (1983), 1243 Friedel pairs  
 Absolute structure parameter: 0.51 (10)

*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 ( $\text{\AA}^2$ )*

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$
Cl2	0.51205 (10)	0.14865 (6)	0.44638 (6)	0.0648 (2)
Cl1	0.44580 (11)	0.75320 (8)	0.58138 (6)	0.0779 (3)
O1	0.3177 (3)	0.44304 (18)	0.42787 (16)	0.0698 (6)
H1	0.2294	0.4174	0.4542	0.105*
N1	0.5869 (3)	0.37691 (19)	0.55437 (15)	0.0522 (6)
H111	0.7029	0.3550	0.5628	0.063*
H112	0.5341	0.3244	0.5198	0.063*
C7	0.4020 (4)	0.5277 (2)	0.4795 (2)	0.0489 (7)
H7	0.3296	0.5422	0.5323	0.059*
C6	0.4249 (3)	0.6368 (2)	0.42750 (19)	0.0488 (6)
C1	0.4496 (3)	0.7418 (2)	0.4678 (2)	0.0546 (7)
C8	0.5901 (4)	0.4868 (2)	0.5047 (2)	0.0537 (7)
H8A	0.6623	0.4771	0.4519	0.064*
H8B	0.6487	0.5443	0.5405	0.064*
C9	0.4922 (4)	0.3739 (2)	0.64202 (18)	0.0579 (7)
H9	0.3639	0.3936	0.6332	0.069*
C5	0.4305 (4)	0.6352 (3)	0.3365 (2)	0.0680 (9)
H5	0.4144	0.5669	0.3069	0.082*
C11	0.5028 (6)	0.2537 (3)	0.6769 (2)	0.0777 (9)
H11A	0.6279	0.2330	0.6854	0.093*
H11B	0.4477	0.2027	0.6355	0.093*
H11C	0.4393	0.2492	0.7319	0.093*
C2	0.4764 (4)	0.8401 (3)	0.4204 (3)	0.0729 (10)
H2	0.4904	0.9091	0.4494	0.088*
C3	0.4824 (5)	0.8360 (4)	0.3318 (3)	0.0920 (13)
H3	0.5018	0.9022	0.2998	0.110*
C4	0.4597 (5)	0.7344 (5)	0.2889 (3)	0.0897 (12)
H4	0.4639	0.7319	0.2279	0.108*
C10	0.5746 (6)	0.4584 (3)	0.7044 (2)	0.0928 (13)

H10A	0.5099	0.4564	0.7591	0.111*
H10B	0.5665	0.5332	0.6797	0.111*
H10C	0.7001	0.4396	0.7144	0.111*

*Atomic displacement parameters (Å<sup>2</sup>)*

	$U^{11}$	$U^{22}$	$U^{33}$	$U^{12}$	$U^{13}$	$U^{23}$
C12	0.0489 (4)	0.0509 (3)	0.0944 (6)	0.0007 (3)	0.0009 (4)	-0.0054 (4)
C11	0.0692 (5)	0.0819 (6)	0.0825 (6)	-0.0038 (5)	-0.0021 (4)	-0.0313 (5)
O1	0.0628 (13)	0.0559 (12)	0.0909 (18)	-0.0136 (10)	-0.0123 (12)	-0.0113 (12)
N1	0.0435 (11)	0.0498 (13)	0.0633 (15)	-0.0017 (10)	-0.0009 (11)	0.0057 (11)
C7	0.0399 (13)	0.0455 (14)	0.0613 (17)	-0.0049 (12)	-0.0016 (12)	-0.0044 (13)
C6	0.0393 (12)	0.0465 (14)	0.0606 (18)	0.0017 (12)	-0.0015 (12)	0.0003 (13)
C1	0.0364 (13)	0.0510 (15)	0.0765 (19)	0.0010 (13)	0.0021 (13)	-0.0009 (15)
C8	0.0438 (14)	0.0483 (15)	0.069 (2)	-0.0023 (13)	0.0006 (13)	0.0073 (13)
C9	0.0513 (15)	0.0647 (18)	0.0577 (17)	-0.0011 (16)	0.0030 (15)	0.0076 (13)
C5	0.0671 (19)	0.080 (2)	0.057 (2)	0.0136 (19)	0.0015 (15)	0.0010 (17)
C11	0.087 (2)	0.074 (2)	0.071 (2)	-0.006 (3)	0.003 (2)	0.0193 (17)
C2	0.0455 (15)	0.0489 (16)	0.124 (3)	-0.0015 (15)	0.007 (2)	0.0108 (18)
C3	0.059 (2)	0.082 (3)	0.135 (4)	0.012 (2)	0.010 (2)	0.050 (3)
C4	0.076 (3)	0.125 (3)	0.068 (2)	0.020 (3)	0.0058 (19)	0.039 (2)
C10	0.123 (3)	0.086 (3)	0.069 (2)	-0.011 (3)	-0.002 (2)	-0.012 (2)

*Geometric parameters (Å, °)*

C11—C1	1.736 (3)	C9—C11	1.514 (4)
O1—C7	1.413 (3)	C9—H9	0.9800
O1—H1	0.8200	C5—C4	1.391 (5)
N1—C8	1.499 (3)	C5—H5	0.9300
N1—C9	1.506 (3)	C11—H11A	0.9600
N1—H111	0.9000	C11—H11B	0.9600
N1—H112	0.9000	C11—H11C	0.9600
C7—C8	1.513 (4)	C2—C3	1.352 (6)
C7—C6	1.518 (4)	C2—H2	0.9300
C7—H7	0.9800	C3—C4	1.373 (7)
C6—C5	1.387 (4)	C3—H3	0.9300
C6—C1	1.393 (4)	C4—H4	0.9300
C1—C2	1.378 (5)	C10—H10A	0.9600
C8—H8A	0.9700	C10—H10B	0.9600
C8—H8B	0.9700	C10—H10C	0.9600
C9—C10	1.503 (5)		
C7—O1—H1	109.5	C10—C9—H9	108.5
C8—N1—C9	118.4 (2)	N1—C9—H9	108.5
C8—N1—H111	107.7	C11—C9—H9	108.5
C9—N1—H111	107.7	C6—C5—C4	121.0 (4)
C8—N1—H112	107.7	C6—C5—H5	119.5
C9—N1—H112	107.7	C4—C5—H5	119.5

H111—N1—H112	107.1	C9—C11—H11A	109.5
O1—C7—C8	108.5 (2)	C9—C11—H11B	109.5
O1—C7—C6	110.8 (2)	H11A—C11—H11B	109.5
C8—C7—C6	107.5 (2)	C9—C11—H11C	109.5
O1—C7—H7	110.0	H11A—C11—H11C	109.5
C8—C7—H7	110.0	H11B—C11—H11C	109.5
C6—C7—H7	110.0	C3—C2—C1	119.9 (3)
C5—C6—C1	116.7 (3)	C3—C2—H2	120.1
C5—C6—C7	120.9 (3)	C1—C2—H2	120.1
C1—C6—C7	122.4 (2)	C2—C3—C4	120.2 (3)
C2—C1—C6	122.2 (3)	C2—C3—H3	119.9
C2—C1—C11	117.4 (3)	C4—C3—H3	119.9
C6—C1—C11	120.4 (2)	C3—C4—C5	120.1 (4)
N1—C8—C7	112.9 (2)	C3—C4—H4	120.0
N1—C8—H8A	109.0	C5—C4—H4	120.0
C7—C8—H8A	109.0	C9—C10—H10A	109.5
N1—C8—H8B	109.0	C9—C10—H10B	109.5
C7—C8—H8B	109.0	H10A—C10—H10B	109.5
H8A—C8—H8B	107.8	C9—C10—H10C	109.5
C10—C9—N1	111.1 (3)	H10A—C10—H10C	109.5
C10—C9—C11	112.1 (3)	H10B—C10—H10C	109.5
N1—C9—C11	108.0 (2)		
O1—C7—C6—C5	-23.6 (4)	C6—C7—C8—N1	-178.2 (2)
C8—C7—C6—C5	94.8 (3)	C8—N1—C9—C10	-58.5 (3)
O1—C7—C6—C1	159.4 (2)	C8—N1—C9—C11	178.2 (3)
C8—C7—C6—C1	-82.2 (3)	C1—C6—C5—C4	0.1 (4)
C5—C6—C1—C2	0.6 (4)	C7—C6—C5—C4	-177.1 (3)
C7—C6—C1—C2	177.8 (2)	C6—C1—C2—C3	-1.1 (4)
C5—C6—C1—C11	-179.8 (2)	C11—C1—C2—C3	179.4 (3)
C7—C6—C1—C11	-2.7 (3)	C1—C2—C3—C4	0.7 (5)
C9—N1—C8—C7	-62.6 (3)	C2—C3—C4—C5	0.0 (6)
O1—C7—C8—N1	-58.4 (3)	C6—C5—C4—C3	-0.5 (5)

Hydrogen-bond geometry (Å, °)

<i>D</i> —H $\cdots$ <i>A</i>	<i>D</i> —H	H $\cdots$ <i>A</i>	<i>D</i> $\cdots$ <i>A</i>	<i>D</i> —H $\cdots$ <i>A</i>
N1—H112 $\cdots$ Cl2	0.90	2.36	3.199 (2)	156
O1—H1 $\cdots$ Cl2 <sup>i</sup>	0.82	2.33	3.143 (2)	169
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Symmetry codes: (i)  $x-1/2, -y+1/2, -z+1$ ; (ii)  $x+1/2, -y+1/2, -z+1$ .