

C(4B)	0.578 (1)	-0.065 (1)	0.3746 (9)	0.078 (7)
C(5B)	0.686 (1)	0.058 (1)	0.3743 (9)	0.072 (6)
C(6B)	0.682 (1)	0.063 (1)	0.259 (1)	0.075 (6)
C(7B)	0.797 (1)	0.172 (1)	0.491 (1)	0.089 (7)
N(8B)	0.363 (1)	-0.2833 (9)	0.0471 (8)	0.096 (6)
C(10B)	0.359 (1)	-0.281 (1)	-0.072 (1)	0.099 (7)
C(12B)	0.465 (2)	-0.319 (1)	-0.122 (1)	0.128 (9)
C(9B)	0.908 (1)	0.287 (1)	0.503 (1)	0.100 (8)
N(14B)	1.014 (1)	0.393 (1)	0.624 (1)	0.113 (8)
O(15B)	1.126 (1)	0.493 (1)	0.6241 (8)	0.145 (7)
O(16B)	0.989 (1)	0.380 (1)	0.7152 (9)	0.143 (7)
C(11B)†	0.236 (2)	-0.421 (2)	0.041 (2)	0.130 (9)
C(13B)‡	0.311 (3)	-0.503 (2)	0.047 (2)	0.121 (9)
C(21B)‡	0.288 (5)	-0.417 (4)	0.054 (3)	
C(23B)‡	0.123 (5)	-0.476 (4)	0.065 (4)	

† Occupancy 0.6.

‡ Occupancy 0.4.

Lists of structure factors, anisotropic displacement parameters and H-atom coordinates have been deposited with the IUCr (Reference: PA1096). Copies may be obtained through The Managing Editor, International Union of Crystallography, 5 Abbey Square, Chester CH1 2HU, England.

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Table 2. Selected geometric parameters ( $\text{\AA}$ , °)

C(1A)—C(2A)	1.43 (2)	C(1B)—C(2B)	1.42 (1)
C(1A)—C(6A)	1.39 (1)	C(1B)—C(6B)	1.35 (1)
C(2A)—C(3A)	1.42 (2)	C(2B)—C(3B)	1.41 (2)
C(2A)—N(8A)	1.356 (9)	C(2B)—N(8B)	1.38 (1)
C(3A)—C(4A)	1.36 (1)	C(3B)—C(4B)	1.37 (1)
C(4A)—C(5A)	1.39 (2)	C(4B)—C(5B)	1.40 (2)
C(5A)—C(6A)	1.41 (2)	C(5B)—C(6B)	1.42 (2)
C(5A)—C(7A)	1.46 (1)	C(5B)—C(7B)	1.47 (1)
C(7A)—C(9A)	1.31 (2)	C(7B)—C(9B)	1.29 (2)
N(8A)—C(10A)	1.48 (2)	N(8B)—C(10B)	1.47 (2)
N(8A)—C(11A)	1.45 (1)	N(8B)—C(11B)	1.55 (2)
C(9A)—N(14A)	1.44 (1)	N(8B)—C(21B)	1.47 (1)
C(10A)—C(12A)	1.49 (2)	C(9B)—N(14B)	1.47 (1)
C(11A)—C(13A)	1.50 (2)	C(10B)—C(12B)	1.50 (2)
N(14A)—O(15A)	1.21 (1)	C(11B)—C(13B)	1.49 (4)
N(14A)—O(16A)	1.23 (2)	C(11B)—C(21B)	0.52 (2)
		C(21B)—C(23B)	1.59 (2)
		N(14B)—O(15B)	1.25 (2)
		N(14B)—O(16B)	1.22 (2)
C(2A)—C(1A)—C(6A)	121 (1)	C(2B)—C(1B)—C(6B)	121 (1)
C(1A)—C(2A)—C(3A)	116.4 (7)	C(1B)—C(2B)—C(3B)	117.2 (7)
C(1A)—C(2A)—N(8A)	120 (1)	C(1B)—C(2B)—N(8B)	121 (1)
C(3A)—C(2A)—N(8A)	124 (1)	C(3B)—C(2B)—N(8B)	121.8 (9)
C(2A)—C(3A)—C(4A)	122 (1)	C(2B)—C(3B)—C(4B)	122 (1)
C(3A)—C(4A)—C(5A)	121 (1)	C(3B)—C(4B)—C(5B)	121 (1)
C(4A)—C(5A)—C(6A)	118.6 (7)	C(4B)—C(5B)—C(6B)	117.2 (7)
C(4A)—C(5A)—C(7A)	120 (1)	C(4B)—C(5B)—C(7B)	118 (1)
C(6A)—C(5A)—C(7A)	122 (1)	C(6B)—C(5B)—C(7B)	124.7 (9)
C(1A)—C(6A)—C(5A)	121 (1)	C(1B)—C(6B)—C(5B)	122 (1)
C(5A)—C(7A)—C(9A)	126 (1)	C(5B)—C(7B)—C(9B)	124 (1)
C(2A)—N(8A)—C(10A)	122.5 (9)	C(2B)—N(8B)—C(10B)	122.4 (9)
C(2A)—N(8A)—C(11A)	121 (1)	C(2B)—N(8B)—C(11B)	123 (1)
C(9A)—N(14A)—O(15A)	117 (1)	C(2B)—N(8B)—C(21B)	114.9 (9)
C(9A)—N(14A)—O(16A)	120 (1)	C(9B)—N(14B)—O(15B)	114 (1)
C(10A)—N(8A)—C(11A)	115.9 (7)	C(9B)—N(14B)—O(16B)	121 (1)
N(8A)—C(10A)—C(12A)	112 (1)	C(10B)—N(8B)—C(11B)	114.1 (9)
N(8A)—C(11A)—C(13A)	114 (1)	C(10B)—N(8B)—C(21B)	119.6 (6)
C(7A)—C(9A)—N(14A)	121 (1)	N(8B)—C(10B)—C(12B)	113 (1)
O(15A)—N(14A)—O(16A)	123.4 (8)	N(8B)—C(11B)—C(13B)	104 (2)
		N(8B)—C(21B)—C(23B)	119 (4)
		C(7B)—C(9B)—N(14B)	120 (1)
		O(15B)—N(14B)—O(16B)	125.2 (9)

In the disordered branch, the pairs [C(11B), C(13B)] and [C21(B), C23(B)] were refined alternatively because of correlations. Isotropic displacement parameters for [C21(B), C23(B)] were not refined.

The structure was solved using PATSEE (Xtal; Hall & Stewart, 1990).

We thank G. Commandeur for the synthesis of the title compound.

*Acta Cryst.* (1995), **C51**, 434–436

## 9-(4-Methyl-1-piperazinyl)-9,10-dihydro-9,10-ethanoanthracene Hydrochloride Dihydrate (Trazitilin)

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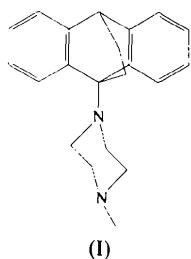
(Received 7 January 1994; accepted 24 June 1994)

## Abstract

The title compound, trazitilin,  $C_{21}H_{24}N_2 \cdot HCl \cdot 2H_2O$ , is an antagonist of histamine H1 receptors. Steric hindrance of the two components of the molecule (the piperazinyl radical and 9,10-dihydro-9,10-ethanoanthracene) causes a slight distortion of substituents and allows the structure very little flexibility.

## Comment

As part of our conformational studies on antihistamine drugs, we investigated the X-ray structure of trazitilin. This compound as a whole is quite rigid and can be used therefore as a basis for the modeling of histamine H1 receptors (Brouant & Barbe, 1992).



Trazitilin crystallizes with two water molecules and one hydrochloride molecule. An intermolecular bonding search reveals that some interactions occur between trazitilin and these molecules. The chloride ion is located at the center of a very flattened tetrahedral complex comprising the two water molecules, the atom N2 from an adjacent cell and one water molecule from a third cell (Fig. 2). The water molecules are themselves connected to atoms of adjacent cells and generate an endless lattice of hydrogen bonds surrounding the molecules of trazitilin. Characteristic contact parameters are reported in Table 3.

The interaction between N2 and the chloride ion is probably the result of the protonation of the N atom; the values of the three angles C1—N2<sup>i</sup>—C21<sup>i</sup> [98.9 (3) $^\circ$ ], C1—N2<sup>i</sup>—C19<sup>i</sup> [103.7 (2) $^\circ$ ] and C1—N2<sup>i</sup>—C18<sup>i</sup> [120.8 (2) $^\circ$ ] (see Fig. 2) lead to this interpretation. No interactions are observed between N1 and other atoms. These results agree with the general observations found for polycyclic compounds substituted by piperazinyl radicals, where the distant N atom is protonated preferentially by acids in solution.

The bonding of the piperazinyl radical to 9,10-dihydro-9,10-ethanoanthracene involves slight dis-

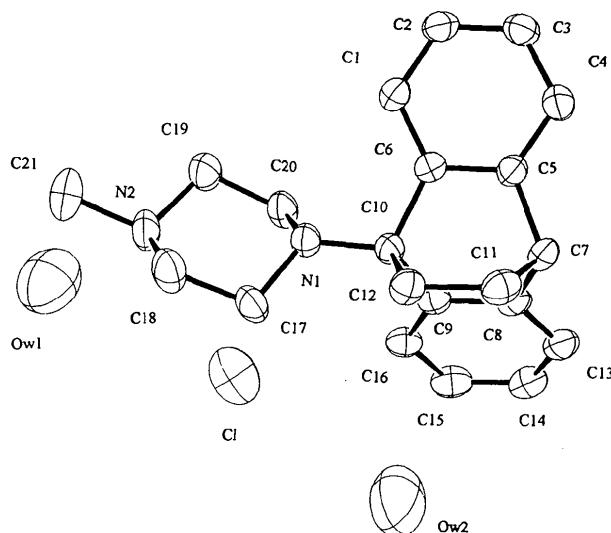


Fig. 1. ORTEPII view (Johnson, 1976) of the title compound. Displacement ellipsoids are drawn at the 50% probability level.

tortions of the latter. These deformations can be described easily in terms of two geometrical features and may be compared to those of similar compounds substituted with smaller components such as methylaminomethyl in benzoctamin (Brouant, Pierrot, Baldy, Soyfer & Barbe, 1984). The ethylene bridge torsion angle C7—C11—C12—C10 is  $-3.7$  (3) $^\circ$  in trazitilin and 1.2 (3) $^\circ$  in benzoctamin, while the angle between the directions C4···C1 and C13···C16 is 4.0 (3) $^\circ$  in trazitilin and 0.5 (2) $^\circ$  in benzoctamin.

Conformational analysis shows that the rotating bond between the polycyclic fragment and the piperazinyl group allows only weak oscillations ( $\pm 7^\circ$  within an energy range of 6 kcal mol $^{-1}$ ; 1 cal = 4.184 J) around the observed value of  $-158.0$  (2) $^\circ$  for C6—C10—N1—C17.

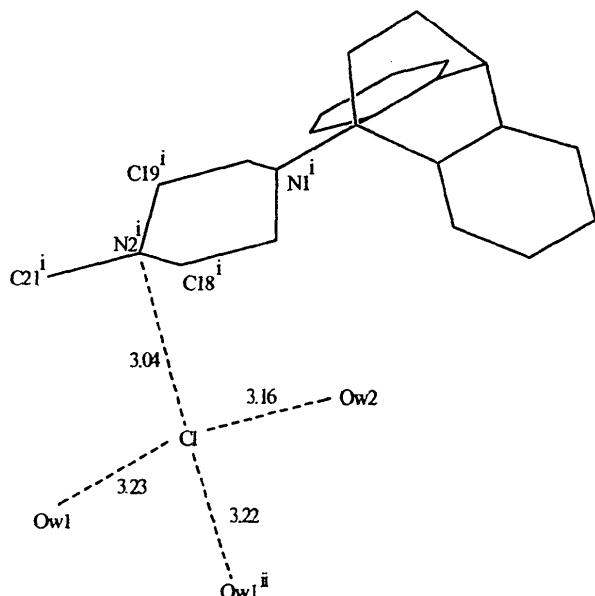
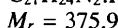
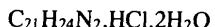


Fig. 2. Intermolecular interactions in the title compound. Symmetry codes: (i)  $x, y, 1 + z$ ; (ii)  $1 - x, 1 - y, 1 - z$ .

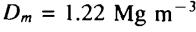
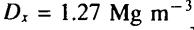
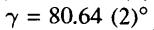
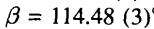
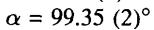
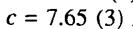
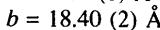
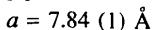
## Experimental

### Crystal data

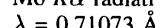


Triclinic

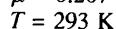
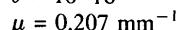
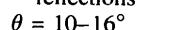
$P\bar{1}$



### Mo K $\alpha$ radiation



Cell parameters from 25 reflections



Orthorhombic

$0.5 \times 0.3 \times 0.3 \text{ mm}$

Transparent

Crystal source: evaporation

of a saturated 50/50%  $\text{H}_2\text{O}/\text{ethanol}$  solution

**Data collection**

Enraf-Nonius CAD-4  
diffractometer  
 $\theta-2\theta$  scans  
Absorption correction:  
none  
3187 measured reflections  
3078 independent reflections  
2064 observed reflections  
[ $F > 3\sigma(F)$ ]

$R_{int} = 0.022$   
 $\theta_{max} = 24^\circ$   
 $h = -8 \rightarrow 8$   
 $k = 0 \rightarrow 21$   
 $l = -8 \rightarrow 8$   
2 standard reflections  
frequency: 60 min  
intensity decay: 2.4%

**Refinement**

Refinement on  $F$   
 $R = 0.051$   
 $wR = 0.054$   
 $S = 0.854$   
2064 reflections  
235 parameters  
H-atom parameters not refined  
Unit weights applied

$(\Delta/\sigma)_{max} = 0.02$   
 $\Delta\rho_{max} = 0.44 (2) e \text{ \AA}^{-3}$   
 $\Delta\rho_{min} = -0.26 (2) e \text{ \AA}^{-3}$   
Extinction correction: none  
Atomic scattering factors  
from *International Tables for X-ray Crystallography*  
(1974, Vol. IV)

C2—C1—C6	119.6 (4)	C6—C10—N1	110.8 (3)
C1—C2—C3	121.2 (3)	C9—C10—C12	104.5 (3)
C2—C3—C4	119.5 (3)	C9—C10—N1	117.8 (3)
C3—C4—C5	119.3 (4)	C12—C10—N1	110.7 (3)
C4—C5—C6	121.3 (3)	C7—C11—C12	109.8 (3)
C4—C5—C7	125.1 (4)	C10—C12—C11	110.2 (3)
C6—C5—C7	113.5 (3)	C8—C13—C14	119.6 (4)
C1—C6—C5	118.9 (3)	C13—C14—C15	120.0 (3)
C1—C6—C10	127.0 (3)	C14—C15—C16	120.7 (3)
C5—C6—C10	114.0 (3)	C9—C16—C15	119.4 (4)
C5—C7—C8	107.5 (3)	C18—C17—N1	109.0 (3)
C5—C7—C11	105.7 (3)	C17—C18—N2	111.4 (3)
C8—C7—C11	107.8 (3)	C20—C19—N2	109.5 (3)
C7—C8—C9	113.9 (3)	C19—C20—N1	108.7 (3)
C7—C8—C13	125.4 (4)	C10—N1—C17	119.3 (3)
C9—C8—C13	120.7 (3)	C10—N1—C20	118.1 (3)
C8—C9—C10	113.3 (3)	C17—N1—C20	108.9 (3)
C8—C9—C16	119.5 (3)	C18—N2—C19	110.3 (3)
C10—C9—C16	127.1 (4)	C18—N2—C21	111.1 (3)
C6—C10—C9	107.4 (3)	C19—N2—C21	111.3 (3)
C6—C10—C12	104.6 (3)		

**Table 1.** Fractional atomic coordinates and equivalent isotropic displacement parameters ( $\text{\AA}^2$ )

$$B_{eq} = (4/3)\sum_i \sum_j \beta_{ij} \mathbf{a}_i \cdot \mathbf{a}_j$$

	$x$	$y$	$z$	$B_{eq}$
C1	0.7057 (5)	0.1129 (2)	0.2381 (5)	2.56 (8)
C2	0.7976 (5)	0.0427 (2)	0.2784 (5)	2.99 (9)
C3	0.7064 (5)	-0.0121 (2)	0.2941 (5)	3.03 (9)
C4	0.5214 (5)	0.0035 (2)	0.2767 (5)	2.86 (8)
C5	0.4295 (4)	0.0735 (2)	0.2369 (5)	2.38 (8)
C6	0.5187 (4)	0.1282 (2)	0.2128 (4)	2.24 (8)
C7	0.2347 (4)	0.1000 (2)	0.2292 (5)	2.69 (8)
C8	0.1224 (4)	0.1310 (2)	0.0402 (5)	2.46 (8)
C9	0.2052 (4)	0.1863 (2)	0.0072 (5)	2.37 (8)
C10	0.3957 (4)	0.2025 (2)	0.1720 (5)	2.35 (8)
C11	0.2599 (5)	0.1647 (2)	0.3911 (5)	3.04 (9)
C12	0.3484 (5)	0.2270 (2)	0.3541 (5)	2.84 (8)
C13	-0.0492 (5)	0.1099 (2)	-0.0936 (5)	3.02 (9)
C14	-0.1386 (5)	0.1431 (2)	-0.2634 (5)	3.38 (9)
C15	-0.0597 (5)	0.1988 (2)	-0.2951 (5)	3.35 (9)
C16	0.1118 (5)	0.2213 (2)	-0.1598 (5)	2.88 (9)
C17	0.4388 (5)	0.3359 (2)	0.1777 (5)	3.38 (9)
C18	0.6030 (6)	0.3794 (2)	0.2262 (6)	4.0 (1)
C19	0.7228 (5)	0.2834 (2)	0.0309 (6)	3.74 (9)
C20	0.5500 (5)	0.2441 (2)	-0.0173 (5)	3.06 (8)
C21	0.8458 (6)	0.4053 (3)	0.1244 (7)	5.7 (1)
N1	0.4987 (4)	0.2571 (2)	0.1482 (4)	2.61 (7)
N2	0.6776 (4)	0.3645 (2)	0.0713 (5)	3.69 (8)
C1	0.4433 (2)	0.41486 (7)	0.6735 (2)	5.55 (3)
Ow1	0.7949 (5)	0.4884 (2)	0.6824 (6)	8.1 (1)
Ow2	0.0506 (6)	0.3676 (2)	0.6113 (7)	10.4 (2)

**Table 2.** Selected geometric parameters ( $\text{\AA}$ ,  $^\circ$ )

C1—C2	1.388 (5)	C10—C12	1.567 (5)
C1—C6	1.384 (5)	C10—N1	1.464 (5)
C2—C3	1.375 (6)	C11—C12	1.550 (6)
C3—C4	1.386 (5)	C13—C14	1.385 (5)
C4—C5	1.383 (5)	C14—C15	1.380 (7)
C5—C6	1.390 (6)	C15—C16	1.393 (5)
C5—C7	1.507 (5)	C17—C18	1.510 (6)
C6—C10	1.536 (4)	C17—N1	1.461 (4)
C7—C8	1.498 (5)	C18—N2	1.497 (6)
C7—C11	1.545 (5)	C19—C20	1.521 (6)
C8—C9	1.399 (6)	C19—N2	1.489 (5)
C8—C13	1.382 (4)	C20—N1	1.456 (5)
C9—C10	1.541 (4)	C21—N2	1.500 (6)
C9—C16	1.387 (5)		

**Table 3.** Intermolecular contact parameters ( $\text{\AA}$ )

Cl—N2 <sup>i</sup>	3.040 (3)	Cl—Ow2	3.160 (5)
Cl—Ow1	3.235 (5)	Ow1—Ow2 <sup>ii</sup>	2.885 (6)
Cl—Ow1 <sup>ii</sup>	3.219 (4)		
N2 <sup>i</sup> —Cl—Ow1	91.1 (1)	Ow1—Cl—Ow1 <sup>ii</sup>	83.3 (1)
N2 <sup>i</sup> —Cl—Ow1 <sup>ii</sup>	164.5 (1)	Ow1—Cl—Ow2	168.6 (1)
N2 <sup>i</sup> —Cl—Ow2	98.9 (1)	Ow1 <sup>ii</sup> —Cl—Ow2	85.6 (1)

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

The H atoms of the water molecules and the N2 proton were not introduced or found by difference Fourier methods. Data collection: CAD-4 software. Data reduction and structure resolution: SDP software package (Frenz, 1978) and direct methods MULTAN80 (Main *et al.*, 1980), completed by Fourier synthesis. H atoms were introduced at idealized positions in the calculation before the last refinement cycle but were not refined. Conformational analysis was performed using SYBYL (TRIPOS, 1993).

Lists of structure factors, anisotropic displacement parameters and H-atom coordinates have been deposited with the IUCr (Reference: PA1108). Copies may be obtained through The Managing Editor, International Union of Crystallography, 5 Abbey Square, Chester CH1 2HU, England.

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