

## 4-Amino-2,2,6,6-tetramethylpiperidin-1-ium chloride monohydrate

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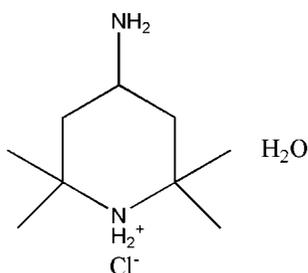
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Key indicators: single-crystal X-ray study;  $T = 113$  K; mean  $\sigma(\text{C}-\text{C}) = 0.002$  Å;  $R$  factor = 0.036;  $wR$  factor = 0.085; data-to-parameter ratio = 18.5.

In the title compound,  $\text{C}_9\text{H}_{21}\text{N}_2^+\cdot\text{Cl}^-\cdot\text{H}_2\text{O}$ , the piperidine ring adopts a chair conformation. In the crystal structure, the components are linked by  $\text{N}-\text{H}\cdots\text{N}$ ,  $\text{O}-\text{H}\cdots\text{Cl}$ ,  $\text{N}-\text{H}\cdots\text{Cl}$  and  $\text{N}-\text{H}\cdots\text{O}$  hydrogen bonds. There are two independent chloride ions, both on crystallographic twofold rotation axes.

### Related literature

For related literature, see: Bojinov &amp; Grabchev (2001).



### Experimental

#### Crystal data

$\text{C}_9\text{H}_{21}\text{N}_2^+\cdot\text{Cl}^-\cdot\text{H}_2\text{O}$   
 $M_r = 210.74$

Orthorhombic,  $Pbcn$   
 $a = 14.1959$  (9) Å

$b = 12.7099$  (7) Å  
 $c = 12.5451$  (6) Å  
 $V = 2263.5$  (2) Å<sup>3</sup>  
 $Z = 8$

Mo  $K\alpha$  radiation  
 $\mu = 0.31$  mm<sup>-1</sup>  
 $T = 113$  (2) K  
 $0.32 \times 0.26 \times 0.20$  mm

#### Data collection

Rigaku Saturn diffractometer  
 Absorption correction: multi-scan  
 (*CrystalClear*; Rigaku/MSC, 2005)  
 $T_{\min} = 0.898$ ,  $T_{\max} = 0.941$

26456 measured reflections  
 2699 independent reflections  
 2672 reflections with  $I > 2\sigma(I)$   
 $R_{\text{int}} = 0.030$

#### Refinement

$R[F^2 > 2\sigma(F^2)] = 0.036$   
 $wR(F^2) = 0.085$   
 $S = 1.15$   
 2699 reflections  
 146 parameters

H atoms treated by a mixture of independent and constrained refinement  
 $\Delta\rho_{\max} = 0.34$  e Å<sup>-3</sup>  
 $\Delta\rho_{\min} = -0.17$  e Å<sup>-3</sup>

**Table 1**

Hydrogen-bond geometry (Å, °).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
$\text{O1}-\text{H1D}\cdots\text{Cl2}$	0.817 (19)	2.44 (2)	3.2509 (11)	174.7 (17)
$\text{O1}-\text{H1C}\cdots\text{Cl1}^{\dagger}$	0.854 (19)	2.244 (19)	3.0971 (11)	177.7 (17)
$\text{N2}-\text{H2C}\cdots\text{Cl2}$	0.882 (18)	2.601 (18)	3.4577 (11)	164.2 (13)
$\text{N1}-\text{H1B}\cdots\text{O1}$	0.894 (17)	2.098 (17)	2.9718 (14)	165.6 (14)
$\text{N1}-\text{H1A}\cdots\text{N2}^{\dagger}$	0.926 (18)	1.935 (18)	2.8602 (14)	176.1 (15)

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

Data collection: *CrystalClear* (Rigaku/MSC, 2005); cell refinement: *CrystalClear*; data reduction: *CrystalClear*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *ORTEP-3* (Farrugia, 1997); software used to prepare material for publication: *CrystalStructure* (Rigaku/MSC, 2005).

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: HB2551).

### References

- Bojinov, V. B. & Grabchev, I. (2001). *Polym. Degrad. Stab.* **74**, 543–550.  
 Farrugia, L. J. (1997). *J. Appl. Cryst.* **30**, 565.  
 Rigaku/MSC (2005). *CrystalClear* and *CrystalStructure*. Rigaku/MSC, The Woodlands, Texas, USA.  
 Sheldrick, G. M. (1997). *SHELXS97* and *SHELXL97*. University of Göttingen, Germany.

**supplementary materials**

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## 4-Amino-2,2,6,6-tetramethylpiperidin-1-ium chloride monohydrate

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### Comment

Hindered amine light stabilizers are one of the most intensively studied classes of stabilizers due to their high photo-stabilization efficiency (Bojinov & Grabchev, 2001). 2,2,6,6-Tetramethylpiperidine-4-amine is an important intermediate of hindered amine light stabilizers.

We report here the crystal structure of the title compound, (I), (Fig. 1), in which the piperidinyll ring adopts a chair conformation. In the crystal, the components are linked by intermolecular N—H $\cdots$ N, O—H $\cdots$ Cl, N—H $\cdots$ Cl and N—H $\cdots$ O hydrogen bonds (Table 1).

### Experimental

The title compound was prepared by dissolving 4-ammonio-2,2,6,6-tetramethylpiperidine in aqueous hydrochloric acid solution. Colourless prisms of (I) were obtained by slow evaporation.

### Refinement

The O- and N-bound H atoms were located in a difference map and their positions and  $U_{\text{iso}}$  values were freely refined. The C-bound atoms were positioned geometrically (C—H = 0.98–0.99 Å), and refined as riding with  $U_{\text{iso}}(\text{H})=1.2U_{\text{eq}}(\text{C})$  or  $1.5U_{\text{eq}}(\text{methyl C})$ .

### Figures



Fig. 1. A view of the molecular structure of (I). Displacement ellipsoids are drawn at the 30% probability level and H atoms are shown as spheres of arbitrary radius. The hydrogen bonds are indicated by dashed lines.

## 4-Amino-2,2,6,6-tetramethylpiperidin-1-ium chloride monohydrate

### Crystal data

$\text{C}_9\text{H}_{21}\text{N}_2^+\cdot\text{Cl}^-\cdot\text{H}_2\text{O}$

$M_r = 210.74$

Orthorhombic, *Pbcn*

$D_x = 1.237 \text{ Mg m}^{-3}$

Melting point: >523 K

Mo  $K\alpha$  radiation

$\lambda = 0.71070 \text{ \AA}$

# supplementary materials

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Hall symbol: -P 2n 2ab

$a = 14.1959 (9) \text{ \AA}$

$b = 12.7099 (7) \text{ \AA}$

$c = 12.5451 (6) \text{ \AA}$

$V = 2263.5 (2) \text{ \AA}^3$

$Z = 8$

$F_{000} = 928$

Cell parameters from 6541 reflections

$\theta = 1.4\text{--}27.9^\circ$

$\mu = 0.31 \text{ mm}^{-1}$

$T = 113 (2) \text{ K}$

Prism, colorless

$0.32 \times 0.26 \times 0.20 \text{ mm}$

## Data collection

Rigaku Saturn  
diffractometer

Radiation source: rotating anode

Monochromator: confocal

$T = 113(2) \text{ K}$

$\omega$  scans

Absorption correction: multi-scan  
(CrystalClear; Rigaku/MSC, 2005)

$T_{\min} = 0.898$ ,  $T_{\max} = 0.941$

26456 measured reflections

2699 independent reflections

2672 reflections with  $I > 2\sigma(I)$

$R_{\text{int}} = 0.030$

$\theta_{\max} = 27.9^\circ$

$\theta_{\min} = 2.2^\circ$

$h = -18 \rightarrow 17$

$k = -16 \rightarrow 16$

$l = -16 \rightarrow 16$

Standard reflections: ?

## Refinement

Refinement on  $F^2$

Least-squares matrix: full

$R[F^2 > 2\sigma(F^2)] = 0.036$

$wR(F^2) = 0.085$

$S = 1.15$

2699 reflections

146 parameters

Primary atom site location: structure-invariant direct methods

Secondary atom site location: difference Fourier map

Hydrogen site location: inferred from neighbouring sites

H atoms treated by a mixture of independent and constrained refinement

$$w = 1/[\sigma^2(F_o^2) + (0.0337P)^2 + 1.133P]$$

where  $P = (F_o^2 + 2F_c^2)/3$

$(\Delta/\sigma)_{\max} = 0.001$

$\Delta\rho_{\max} = 0.34 \text{ e \AA}^{-3}$

$\Delta\rho_{\min} = -0.17 \text{ e \AA}^{-3}$

Extinction correction: SHELXL97 (Sheldrick, 1997),

$$F_c^* = kFc[1 + 0.001xFc^2\lambda^3/\sin(2\theta)]^{-1/4}$$

Extinction coefficient: 0.0104 (8)

## 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  $F^2 > 2\sigma(F^2)$  is used only for calculat-

ing  $R$ -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 ( $\text{\AA}^2$ )

	$x$	$y$	$z$	$U_{\text{iso}}^*/U_{\text{eq}}$
Cl1	0.0000	0.64751 (3)	0.2500	0.02249 (13)
Cl2	0.5000	0.56016 (3)	0.2500	0.01639 (12)
N1	0.25307 (7)	0.33787 (8)	0.40843 (8)	0.0111 (2)
H1A	0.2197 (11)	0.2761 (14)	0.4188 (13)	0.026 (4)*
H1B	0.2920 (11)	0.3283 (12)	0.3533 (13)	0.020 (4)*
N2	0.35317 (8)	0.65104 (8)	0.44971 (9)	0.0143 (2)
H2C	0.3942 (12)	0.6415 (12)	0.3978 (14)	0.020 (4)*
H2D	0.3836 (12)	0.6657 (13)	0.5061 (14)	0.025 (4)*
O1	0.37732 (7)	0.34651 (8)	0.21798 (8)	0.0200 (2)
H1C	0.4126 (13)	0.2927 (15)	0.2264 (14)	0.030*
H1D	0.4113 (13)	0.3976 (15)	0.2268 (14)	0.030*
C1	0.31784 (8)	0.35426 (9)	0.50387 (9)	0.0118 (2)
C2	0.36764 (8)	0.45990 (9)	0.48955 (9)	0.0125 (2)
H2A	0.4132	0.4538	0.4301	0.015*
H2B	0.4038	0.4758	0.5551	0.015*
C3	0.30101 (8)	0.55161 (9)	0.46649 (9)	0.0121 (2)
H3	0.2575	0.5606	0.5286	0.015*
C4	0.24252 (8)	0.52763 (9)	0.36733 (9)	0.0126 (2)
H4A	0.2852	0.5226	0.3051	0.015*
H4B	0.1988	0.5869	0.3544	0.015*
C5	0.18539 (8)	0.42568 (9)	0.37562 (9)	0.0124 (2)
C6	0.38975 (9)	0.26463 (9)	0.50102 (10)	0.0157 (3)
H6A	0.3573	0.1973	0.5111	0.024*
H6B	0.4359	0.2746	0.5582	0.024*
H6C	0.4221	0.2646	0.4320	0.024*
C7	0.26345 (9)	0.34794 (9)	0.60886 (9)	0.0154 (3)
H7A	0.2233	0.4102	0.6163	0.023*
H7B	0.3080	0.3450	0.6685	0.023*
H7C	0.2242	0.2845	0.6091	0.023*
C8	0.14726 (9)	0.39370 (10)	0.26623 (10)	0.0165 (3)
H8A	0.1999	0.3833	0.2168	0.025*
H8B	0.1059	0.4493	0.2390	0.025*
H8C	0.1115	0.3281	0.2728	0.025*
C9	0.10232 (9)	0.43610 (10)	0.45239 (10)	0.0166 (3)
H9A	0.0752	0.3664	0.4659	0.025*
H9B	0.0543	0.4819	0.4208	0.025*
H9C	0.1242	0.4666	0.5197	0.025*

Atomic displacement parameters ( $\text{\AA}^2$ )

	$U^{11}$	$U^{22}$	$U^{33}$	$U^{12}$	$U^{13}$	$U^{23}$
Cl1	0.0233 (2)	0.0148 (2)	0.0293 (3)	0.000	-0.00737 (18)	0.000

## supplementary materials

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C12	0.0156 (2)	0.0194 (2)	0.0141 (2)	0.000	0.00074 (14)	0.000
N1	0.0117 (5)	0.0105 (5)	0.0110 (4)	0.0000 (4)	0.0002 (4)	-0.0006 (3)
N2	0.0153 (5)	0.0109 (5)	0.0166 (5)	-0.0014 (4)	-0.0005 (4)	0.0000 (4)
O1	0.0166 (5)	0.0188 (5)	0.0246 (5)	0.0000 (4)	-0.0012 (4)	-0.0020 (4)
C1	0.0135 (5)	0.0113 (5)	0.0106 (5)	0.0005 (4)	-0.0014 (4)	0.0003 (4)
C2	0.0129 (5)	0.0114 (5)	0.0131 (5)	-0.0007 (4)	-0.0005 (4)	0.0001 (4)
C3	0.0136 (5)	0.0104 (5)	0.0124 (5)	-0.0004 (4)	0.0007 (4)	-0.0001 (4)
C4	0.0141 (5)	0.0111 (5)	0.0128 (5)	0.0006 (4)	-0.0004 (4)	0.0014 (4)
C5	0.0116 (5)	0.0115 (5)	0.0140 (5)	0.0026 (4)	-0.0008 (4)	0.0007 (4)
C6	0.0165 (6)	0.0130 (6)	0.0176 (6)	0.0013 (4)	-0.0020 (4)	0.0009 (4)
C7	0.0188 (6)	0.0158 (6)	0.0115 (5)	-0.0008 (5)	0.0006 (5)	0.0015 (4)
C8	0.0171 (6)	0.0157 (6)	0.0167 (6)	0.0013 (5)	-0.0038 (5)	-0.0015 (5)
C9	0.0130 (5)	0.0167 (6)	0.0200 (6)	0.0009 (4)	0.0022 (5)	-0.0001 (5)

### Geometric parameters ( $\text{\AA}$ , $^\circ$ )

N1—C1	1.5240 (14)	C4—C5	1.5322 (16)
N1—C5	1.5290 (14)	C4—H4A	0.9900
N1—H1A	0.926 (18)	C4—H4B	0.9900
N1—H1B	0.894 (17)	C5—C9	1.5283 (16)
N2—C3	1.4798 (15)	C5—C8	1.5302 (16)
N2—H2C	0.882 (18)	C6—H6A	0.9800
N2—H2D	0.850 (18)	C6—H6B	0.9800
O1—H1C	0.854 (19)	C6—H6C	0.9800
O1—H1D	0.817 (19)	C7—H7A	0.9800
C1—C2	1.5281 (15)	C7—H7B	0.9800
C1—C7	1.5287 (16)	C7—H7C	0.9800
C1—C6	1.5301 (16)	C8—H8A	0.9800
C2—C3	1.5288 (16)	C8—H8B	0.9800
C2—H2A	0.9900	C8—H8C	0.9800
C2—H2B	0.9900	C9—H9A	0.9800
C3—C4	1.5264 (16)	C9—H9B	0.9800
C3—H3	1.0000	C9—H9C	0.9800
C1—N1—C5	119.39 (9)	C5—C4—H4B	108.8
C1—N1—H1A	108.3 (10)	H4A—C4—H4B	107.7
C5—N1—H1A	109.6 (10)	C9—C5—N1	112.24 (9)
C1—N1—H1B	104.7 (10)	C9—C5—C8	108.37 (10)
C5—N1—H1B	106.2 (10)	N1—C5—C8	105.65 (9)
H1A—N1—H1B	108.1 (14)	C9—C5—C4	112.21 (10)
C3—N2—H2C	108.6 (10)	N1—C5—C4	107.64 (9)
C3—N2—H2D	108.9 (12)	C8—C5—C4	110.55 (9)
H2C—N2—H2D	108.0 (15)	C1—C6—H6A	109.5
H1C—O1—H1D	105.9 (18)	C1—C6—H6B	109.5
N1—C1—C2	107.87 (9)	H6A—C6—H6B	109.5
N1—C1—C7	111.41 (9)	C1—C6—H6C	109.5
C2—C1—C7	112.41 (9)	H6A—C6—H6C	109.5
N1—C1—C6	106.40 (9)	H6B—C6—H6C	109.5
C2—C1—C6	110.04 (10)	C1—C7—H7A	109.5
C7—C1—C6	108.54 (9)	C1—C7—H7B	109.5

C1—C2—C3	113.95 (9)	H7A—C7—H7B	109.5
C1—C2—H2A	108.8	C1—C7—H7C	109.5
C3—C2—H2A	108.8	H7A—C7—H7C	109.5
C1—C2—H2B	108.8	H7B—C7—H7C	109.5
C3—C2—H2B	108.8	C5—C8—H8A	109.5
H2A—C2—H2B	107.7	C5—C8—H8B	109.5
N2—C3—C4	109.07 (9)	H8A—C8—H8B	109.5
N2—C3—C2	111.62 (9)	C5—C8—H8C	109.5
C4—C3—C2	109.78 (9)	H8A—C8—H8C	109.5
N2—C3—H3	108.8	H8B—C8—H8C	109.5
C4—C3—H3	108.8	C5—C9—H9A	109.5
C2—C3—H3	108.8	C5—C9—H9B	109.5
C3—C4—C5	113.67 (9)	H9A—C9—H9B	109.5
C3—C4—H4A	108.8	C5—C9—H9C	109.5
C5—C4—H4A	108.8	H9A—C9—H9C	109.5
C3—C4—H4B	108.8	H9B—C9—H9C	109.5
C5—N1—C1—C2	-50.01 (13)	N2—C3—C4—C5	179.86 (9)
C5—N1—C1—C7	73.82 (12)	C2—C3—C4—C5	57.27 (13)
C5—N1—C1—C6	-168.06 (9)	C1—N1—C5—C9	-73.64 (13)
N1—C1—C2—C3	51.43 (12)	C1—N1—C5—C8	168.44 (10)
C7—C1—C2—C3	-71.78 (13)	C1—N1—C5—C4	50.30 (13)
C6—C1—C2—C3	167.12 (9)	C3—C4—C5—C9	71.93 (12)
C1—C2—C3—N2	-178.05 (9)	C3—C4—C5—N1	-52.03 (12)
C1—C2—C3—C4	-56.98 (13)	C3—C4—C5—C8	-166.96 (10)

Hydrogen-bond geometry ( $\text{\AA}$ ,  $^\circ$ )

<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>
O1—H1D $\cdots$ Cl2	0.817 (19)	2.44 (2)	3.2509 (11)	174.7 (17)
O1—H1C $\cdots$ Cl1 <sup>i</sup>	0.854 (19)	2.244 (19)	3.0971 (11)	177.7 (17)
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N1—H1B $\cdots$ O1	0.894 (17)	2.098 (17)	2.9718 (14)	165.6 (14)
N1—H1A $\cdots$ N2 <sup>i</sup>	0.926 (18)	1.935 (18)	2.8602 (14)	176.1 (15)

Symmetry codes: (i)  $-x+1/2, y-1/2, z$ .

Fig. 1

