organic papers

Acta Crystallographica Section E Structure Reports Online

ISSN 1600-5368

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Key indicators

Single-crystal X-ray study T = 295 KMean $\sigma(\text{C}-\text{C}) = 0.004 \text{ Å}$ R factor = 0.051 wR factor = 0.092 Data-to-parameter ratio = 22.5

For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e.

(3a*R*,8b*R*)-(+)-1-Methyl-1,2,3,3a,4,8bhexahydropyrrolo[2',3':3,4]cyclopenta-[1,2-*b*]pyridine-1,5-diium dichloride monohydrate

The title compound, $C_{11}H_{16}N_2^{2+}\cdot 2Cl^-\cdot H_2O$, is a novel conformationally constrained tricyclic analogue of nicotine, in which the pyridine and pyrrolidine rings of the latter are bridged by a CH_2 group. As result of this bridge, these two rings are no longer approximately perpendicular as in various salts of nicotine.

Comment

The alkaloid (S)-(-)-nicotine, (II), is a mimic of the important neurotransmitter acetylcholine and interacts with a wide range of biological binding sites, including those in acetylcholinesterases and nicotinic acetylcholine receptors (nAChR) (Schmitt & Bencherif, 2000). Such receptors have been implicated in a number of cognitive and learning processes, making them a promising target for the treatment of neurodegenerative diseases, such as Alzheimer's or Parkinson's diseases (Meyer *et al.*, 2000; Levin & Simon, 1998). In this context, and in view of the alkaloid epibatidine (Spande *et al.*, 1992), conformationally constrained nicotine analogues became of interest because restrictions in conformation may potentially tune potency and selectivity, while reducing toxicity in comparison with nicotine (Glassco *et al.*, 1993).



The title compound, (I), is such a derivative, having the *N*-methylpyrrolidine and pyridine units of nicotine fixed relative to each other *via* a bridging CH_2 group (Ullrich *et al.*, 2002). As the compound was synthesized in a non-enantioselective route and optically resolved with a chiral auxiliary (Binder & Pyerin, 2005), it was necessary to determine its absolute configuration. The result is shown in Fig. 1. The chiral centre at position 2 of the pyrrolidine ring in the (+)-enantiomer of (I) has a configuration opposite to that of the analogous chiral centre in (II).

Whereas covalent bond lengths are normal (Table 1; Allen *et al.*, 1999), bond angles in the central five-membered ring C1/C2/C6/C7/C11 (e.g. C1-C2-C6 and C1-C11-C7) indicate moderate strain and the expected rigidity of the whole molecule. The central five-membered ring is almost flat and coplanar with the pyridine ring (r.m.s. deviation of both rings

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Figure 1

A view of the asymmetric unit of (I), with displacement ellipsoids drawn at the 30% probability level. Dashed lines indicate hydrogen bonds.



Figure 2

A packing diagram for (I), viewed down the b axis, with hydrogen bonds shown as dashed lines. C-bound H atoms have been omitted for clarity.

from a common least-squares plane is 0.023 Å). The pyrrolidine ring has an envelope conformation, with atom C9 deviating by 0.592 (4) Å from the least-squares plane of the other four ring atoms.

Each N atom is protonated and donates a comparatively short hydrogen bond to a neighbouring Cl⁻ ion (Table 2). The water molecule is hydrogen bonded to Cl⁻ ions of two adjacent $C_{11}H_{16}N_2^{2+}\cdot 2Cl^-$ units and links them into continuous hydrogen-bonded chains parallel to the *a* axis (Fig. 2).

Compared with typical compounds of nicotine, the molecule of (I) has an atypical shape, as a result of the bridging carbon C11. This can be demonstrated by the torsion angle C1-C2-C6-H6, which is -129° in (I), but in other nicotine compounds is either near 0° [-10° and 38° in nicotinium tetrachlorocuprate(II) (Choi et al., 2002) and 3° in nicotinium iodide (Barlow et al., 1986)] or near 180° (167° in nicotinium salicylate; Kim & Jeffrey, 1971).

Experimental

The synthesis of (I) was recently described by Binder & Pyerin (2005). Crystals for X-ray analysis in the form of thin colourless prisms were obtained by recrystallization from ethanol-H₂O (15:1) by slow evaporation at room temperature. $\left[\alpha\right]_{D}^{20} 27.5^{\circ}$ (c = 0.42, methanol).

Z = 4

 $D_x = 1.358 \text{ Mg m}^{-3}$

Mo $K\alpha$ radiation

Prism, colourless

 $0.55 \times 0.07 \times 0.03 \text{ mm}$

17008 measured reflections

3554 independent reflections

2829 reflections with $I > 2\sigma(I)$

 $\mu = 0.48 \text{ mm}^{-1}$

T = 295 (2) K

 $R_{\rm int} = 0.043$

 $\theta_{\rm max} = 30.0^{\circ}$

Crystal data

 $C_{11}H_{16}N_2^{2+}\cdot 2Cl^-\cdot H_2O$ $M_r = 265.17$ Orthorhombic, $P2_12_12_1$ a = 6.853 (3) Å b = 8.789 (4) Å c = 21.537 (9) Å V = 1297.2 (10) Å³

Data collection

Bruker SMART CCD area-detector diffractometer ω scans Absorption correction: multi-scan (SADABS; Bruker, 2003) $T_{\rm min} = 0.92, T_{\rm max} = 0.99$

Refinement

Table 1

T.L.L. 0

| Refinement on F^2 | $w = 1/[\sigma^2(F_0^2) + (0.0323P)^2]$ |
|---------------------------------|--|
| $R[F^2 > 2\sigma(F^2)] = 0.051$ | + 0.2916P] |
| $vR(F^2) = 0.092$ | where $P = (F_0^2 + 2F_c^2)/3$ |
| S = 1.12 | $(\Delta/\sigma)_{\rm max} < 0.001$ |
| 554 reflections | $\Delta \rho_{\rm max} = 0.23 \ {\rm e} \ {\rm \AA}^{-3}$ |
| 58 parameters | $\Delta \rho_{\rm min} = -0.20 \text{ e } \text{\AA}^{-3}$ |
| I atoms treated by a mixture of | Absolute structure: Flack (1983), |
| independent and constrained | with 1449 Friedel pairs |
| refinement | Flack parameter: 0.03 (7) |

Selected geometric parameters (Å, °).

| N1-C1 | 1.346 (3) | C1-C11 | 1.474 (4) |
|----------------------------|------------------------|----------------------------|------------------------|
| N1-C5 | 1.336 (3) | C2-C6 | 1.497 (3) |
| N2-C6 | 1.523 (3) | C6-C7 | 1.556 (3) |
| N2-C9 | 1.501 (3) | C7-C8 | 1.520 (4) |
| N2-C10 | 1.493 (3) | C8-C9 | 1.507 (4) |
| C1-C2 | 1.370 (3) | C7-C11 | 1.553 (3) |
| C1-C2-C6 C3-C2-C6 | 110.0 (2) 129.6 (2) | C1-C11-C7 | 103.8 (2) |
| C1-C2-C6-C7 C1-C2-C6-N2 | -1.9 (2) 112.2 (2) | C2-C6-C7-C8 C6-C7-C8-C9 | 124.5 (2) -29.4 (3) |

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

| $D - H \cdots A$ | D-H | $H \cdot \cdot \cdot A$ | $D \cdots A$ | $D - \mathbf{H} \cdot \cdot \cdot A$ |
|---|--|--|--|--|
| $N1 - H1N \cdots Cl1$ $N2 - H2N \cdots Cl2$ $O1 - H1OA \cdots Cl1$ $O1 - H1OB \cdots Cl2^{i}$ | 0.91 (3) 0.95 (3) 0.99 (4) 0.96 (4) | 2.09 (3) 2.13 (3) 2.39 (4) 2.31 (4) | 3.000 (2) 3.075 (2) 3.351 (3) 3.240 (3) | 174 (2) 174 (2) 164 (3) 163 (3) |
| | | | | |

Symmetry code: (i) x - 1, y, z.

All C-bound H atoms were placed in calculated positions (C-H =0.93-0.98 Å) and thereafter treated as riding. A torsional parameter was refined for the methyl group. The positions of N- and O-bound H atoms were refined freely (distances are in Table 2). For all H atoms, $U_{iso}(H) = 1.2U_{eq}(C,N,O)$ or $U_{iso}(H) = 1.5U_{eq}(C_{methyl})$.

Data collection: *SMART* (Bruker, 2003); cell refinement: *SAINT* (Bruker, 2003); data reduction: *SAINT*, *SADABS* and *XPREP* (Bruker, 2003); program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *SHELXTL* (Bruker, 2003); software used to prepare material for publication: *SHELXTL*.

The authors thank Gerhard Denk, René Wissiack, and Heinz A. Krebs for their assistance in the synthesis of key compounds.

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