

Refinement

Refinement on F^2
 $R[F^2 > 2\sigma(F^2)] = 0.038$
 $wR(F^2) = 0.104$
 $S = 0.905$
3274 reflections
238 parameters
H-atom parameters
constrained
 $w = 1/[\sigma^2(F_o^2) + (0.0574P)^2]$
where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\text{max}} < 0.001$

$\Delta\rho_{\text{max}} = 0.182 \text{ e } \text{\AA}^{-3}$
 $\Delta\rho_{\text{min}} = -0.172 \text{ e } \text{\AA}^{-3}$
Extinction correction:
SHELXL97 (Sheldrick, 1997a)
Extinction coefficient:
0.0069 (10)
Scattering factors from
International Tables for Crystallography (Vol. C)

Table 1. Selected geometric parameters (\AA , $^\circ$)

O1—C12	1.359 (2)	C11—C21	1.497 (2)
O2—C22	1.356 (2)	C31—C41	1.480 (2)
O1—C12—C11	118.11 (16)	O2—C22—C23	121.90 (17)
O1—C12—C13	121.69 (18)	C33—N34—C35	116.02 (18)
O2—C22—C21	117.91 (16)	C43—N44—C45	116.02 (17)
C16—C11—C21—C26	−124.4 (2)	C36—C31—C41—C42	21.2 (3)
C12—C11—C21—C26	55.6 (2)	C32—C31—C41—C42	−162.8 (2)
C16—C11—C21—C22	55.2 (2)	C36—C31—C41—C46	−156.7 (2)
C12—C11—C21—C22	−124.8 (2)	C32—C31—C41—C46	19.3 (3)

Table 2. Hydrogen-bonding geometry (\AA , $^\circ$)

$D—H \cdots A$	$D—H$	$H \cdots A$	$D \cdots A$	$D—H \cdots A$
O1—H1—N34	0.82	1.98	2.793 (3)	172
O2—H2—N44 ^t	0.82	1.96	2.750 (3)	161

Symmetry code: (i) $\frac{1}{2} - x, y - \frac{3}{2}, \frac{1}{2} - z$.

Compound (I) crystallized in the monoclinic system. From the systematic absences, the space group was $P2_1/n$. H atoms were treated as riding atoms (C—H 0.93 and O—H 0.82 \AA).

Data collection: *CAD-4-PC Software* (Enraf–Nonius, 1992). Cell refinement: *SET4* and *CELDIM* (*CAD-4-PC Software*; Enraf–Nonius, 1992). Data reduction: *DATRD2* in *NRCVAX96* (Gabe *et al.*, 1989). Program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997b). Program(s) used to refine structure: *NRCVAX96* and *SHELXL97* (Sheldrick, 1997a). Molecular graphics: *NRCVAX96*, *ORTEPII* (Johnson, 1976) and *PLATON* (Spek, 1998). Software used to prepare material for publication: *NRCVAX96*, *SHELXL97* and *WordPerfect macro PREP8* (Ferguson, 1998).

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Supplementary data for this paper are available from the IUCr electronic archives (Reference: SK1245). Services for accessing these data are described at the back of the journal.

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Paroxetine hydrochloride hemihydrate

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Abstract

The structure of the title compound [systematic name: (3S-trans)-3-[(1,3-benzodioxol-5-yloxy)methyl]-4-(4-fluorophenyl)piperidinium chloride hemihydrate, $C_{19}H_{21}FNO_3^{\pm}\cdot Cl^- \cdot 0.5H_2O$], comprises two piperidinium cations (*A* and *B*), two chloride anions, and one water molecule in the asymmetric unit. A piperidinium proton of cation *A* is hydrogen bonded to the water molecule. The second proton points in the direction of one chloride anion, as does a piperidinium proton of cation *B*. One water proton points in the direction of another chloride anion.

Comment

Paroxetine hydrochloride hemihydrate is a serotonin (5-HT) uptake inhibitor and is of therapeutic use as an antidepressant (Barnes *et al.*, 1988). There is increased interest in the use of paroxetine as an alternative to the benzodiazepines in the treatment of generalized anxiety disorder (Rocca *et al.*, 1997). Paroxetine hydrochloride hemihydrate, (I), can be dehydrated, although the hemihydrate is more stable (Buxton *et al.*, 1988). Recently, a variety of ‘anhydrides’ have been reported

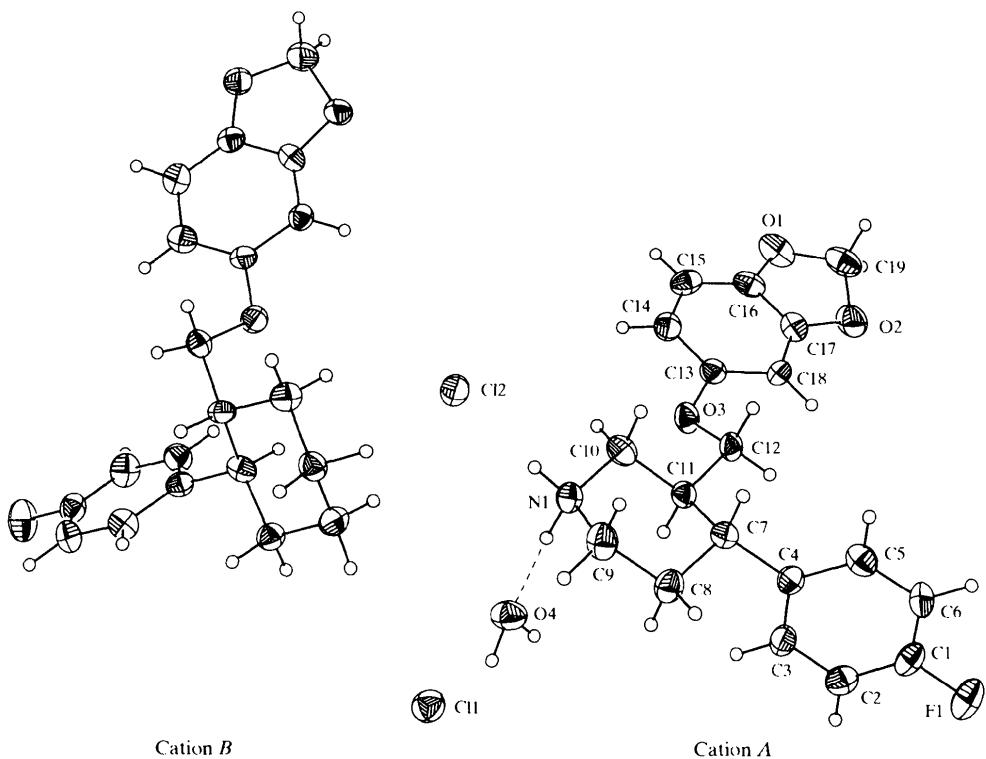
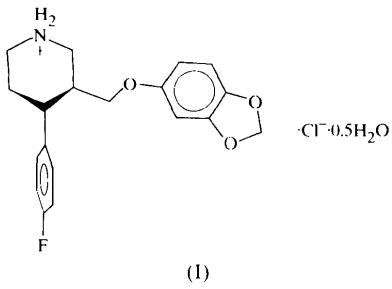


Fig. 1. The structure of paroxetine hydrochloride hemihydrate with 50% probability displacement ellipsoids, except for H atoms, which are drawn as small circles.

(Jacewicz & Ward, 1996), but these have not been characterized structurally. We report here the first crystal structure of a paroxetine hydrochloride compound.



The structure of the hemihydrate (I) confirms the basic composition and stereochemistry of paroxetine, as derived by spectroscopic and chemical means. The water molecule is bound to one of the paroxetine molecules through an N—H \cdots O hydrogen bond 2.772(6) Å in length. That paroxetine hydrochloride hemihydrate is readily dehydrated is thus not surprising.

Experimental

The title compound was obtained from Schein Pharmaceutical, Danbury, Connecticut, USA.

Crystal data

$C_{19}H_{21}FNO_3\cdot Cl^- \cdot 0.5H_2O$	Mo $K\alpha$ radiation
$M_r = 374.83$	$\lambda = 0.71073 \text{ \AA}$
Monoclinic	Cell parameters from 1828 reflections
$P2_1$	$\theta = 1-25^\circ$
$a = 12.8953(17) \text{ \AA}$	$\mu = 0.244 \text{ mm}^{-1}$
$b = 10.0927(11) \text{ \AA}$	$T = 153(2) \text{ K}$
$c = 14.413(2) \text{ \AA}$	Needle
$\beta = 106.638(12)^\circ$	$0.29 \times 0.076 \times 0.044 \text{ mm}$
$V = 1797.2(4) \text{ \AA}^3$	Colourless
$Z = 4$	
$D_x = 1.385 \text{ Mg m}^{-3}$	
D_m not measured	

Data collection

Bruker Smart 1000 CCD diffractometer	3802 reflections with $I > 2\sigma(I)$
ω scans	$R_{\text{int}} = 0.065$
Absorption correction: numerical, face indexed (Sheldrick, 1997)	$\theta_{\text{max}} = 25^\circ$
$T_{\text{min}} = 0.954$, $T_{\text{max}} = 0.991$	$h = -15 \rightarrow 15$
10 872 measured reflections	$k = -11 \rightarrow 12$
6093 independent reflections	$l = -17 \rightarrow 15$
	Intensity decay: <2%

Refinement

Refinement on F^2	$(\Delta/\sigma)_{\text{max}} = 0.014$
$R(F) = 0.044$	$\Delta\rho_{\text{max}} = 0.24 \text{ e \AA}^{-3}$
$wR(F^2) = 0.089$	$\Delta\rho_{\text{min}} = -0.20 \text{ e \AA}^{-3}$

$S = 0.78$
 6093 reflections
 484 parameters
 H atoms treated by a mixture of independent and constrained refinement

Extinction correction: none
 Scattering factors from *International Tables for Crystallography* (Vol. C)
 Absolute structure: Flack (1983)
 Flack parameter = -0.06 (6)

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Table 1. Selected geometric parameters (\AA , $^\circ$)

C1A—F1A	1.365 (5)	C16A—C17A	1.368 (6)
C1B—F1B	1.366 (5)	C16A—O1A	1.384 (5)
C4A—C7A	1.518 (6)	C16B—C17B	1.383 (6)
C4B—C7B	1.518 (5)	C16B—O1B	1.392 (5)
C7A—C11A	1.537 (5)	C17A—O2A	1.375 (5)
C7A—C8A	1.541 (5)	C17B—O2B	1.370 (4)
C7B—C8B	1.538 (5)	C19A—O1A	1.417 (6)
C7B—C11B	1.539 (5)	C19A—O2A	1.445 (5)
C8A—C9A	1.517 (6)	C19B—O2B	1.418 (5)
C8B—C9B	1.516 (5)	C19B—O1B	1.452 (5)
C9A—N1A	1.496 (5)	C11···H1O4 ^a	2.35 (5)
C9B—N1B	1.490 (5)	C12···H2N2	1.94 (5)
C10A—N1A	1.475 (5)	C12···H1N1	2.18 (4)
C10A—C11A	1.523 (6)	N1A—H1N1	0.89 (4)
C10B—N1B	1.483 (5)	N1A—H2N1	1.12 (5)
C10B—C11B	1.542 (5)	N1B—H1N2	0.84 (4)
C11A—C12A	1.524 (5)	N1B—H2N2	1.22 (5)
C11B—C12B	1.501 (5)	O4—H1O4	0.79 (5)
C12A—O3A	1.428 (4)	O4—H2O4	0.99 (7)
C12B—O3B	1.428 (4)	O4···H2N1	1.69 (5)
C13A—O3A	1.381 (4)	O4···N1A	2.772 (6)
C13B—O3B	1.371 (4)		
C4A—C7A—C11A	114.5 (4)	C16A—C17A—O2A	110.0 (4)
C4A—C7A—C8A	111.3 (3)	C18A—C17A—O2A	127.1 (4)
C4B—C7B—C8B	112.5 (3)	O2B—C17B—C18B	128.3 (4)
C4B—C7B—C11B	109.7 (3)	O2B—C17B—C16B	109.9 (3)
O3A—C12A—C11A	105.8 (3)	O1A—C19A—O2A	107.2 (4)
O3B—C12B—C11B	108.7 (3)	O2B—C19B—O1B	107.0 (3)
O3A—C13A—C14A	115.7 (4)	H1N1—N1A—H2N1	112 (4)
O3A—C13A—C18A	122.8 (4)	H1N2—N1B—H2N2	114 (4)
O3B—C13B—C14B	123.3 (4)	H1O4—O4—H2O4	94 (5)
O3B—C13B—C18B	115.2 (4)	N1A—H1N1···C12	174 (4)
C15A—C16A—O1A	128.4 (4)	N1A—H2N1···O4	159 (4)
C17A—C16A—O1A	109.7 (4)	N1B—H2N2···C12	152 (3)
C15B—C16B—O1B	128.7 (4)	O4—H1O4···C11 ^b	164 (4)
C17B—C16B—O1B	109.2 (4)	O4—H2O4···C11	143 (6)

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

The positional and isotropic displacement parameters of the H atoms on the piperidinium N atoms and on the water molecule were refined. Other H atoms were placed at calculated positions and were refined with a riding model. All non-H atoms were refined anisotropically.

Data collection: SMART (Bruker, 1997). Cell refinement: SMART. Data reduction: SAINT-Plus (Bruker, 1997). Program(s) used to solve structure: SHELXS97 (Sheldrick, 1990). Program(s) used to refine structure: SHELXTL/PC (Sheldrick, 1997). Molecular graphics: SHELXTL/PC. Software used to prepare material for publication: SHELXTL/PC.

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Supplementary data for this paper are available from the IUCr electronic archives (Reference: FR1162). Services for accessing these data are described at the back of the journal.

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Fromm's methylphenyldithiodimethylketuret

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Abstract

The 'keturet' obtained by reacting 1-methyl-1-phenyldithiobiuret with acetone is shown to be 6,6-dimethyl-2-methylanilino-5,6-dihydro-4*H*-1,3,5-thiadiazine-4-thione, $C_{12}H_{15}N_3S_2$. The molecule contains a six-membered thiadiazine ring with a distorted half-chair conformation, rather than a previously proposed four-membered ring structure. Molecular dimensions are normal, and the molecules (two per asymmetric unit) are linked by two N—H···S hydrogen bonds, one fairly strong [H···S 2.39 (3) \AA] and one weaker [H···S 2.65 (4) \AA].

Comment

Condensation of 1-aryldithiobiurets with ketones, under the influence of dry hydrogen chloride, gives compounds called 'keturets' (Fromm, 1893), some of which can be formulated as hexahydrotriazine derivatives (Fairfull & Peak, 1955). Reaction of 1-methyl-1-phenyldithiobiuret, (I), with acetone gives a keturet which cannot be a hexahydrotriazine (since the disubstitution on N1