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5-(4-Fluorophenyl)-1,8-dimethyl-2-(*p*-toluoylaminoethyl)-2,3-dihydro-1*H*-1,4-benzodiazepine Monohydrate

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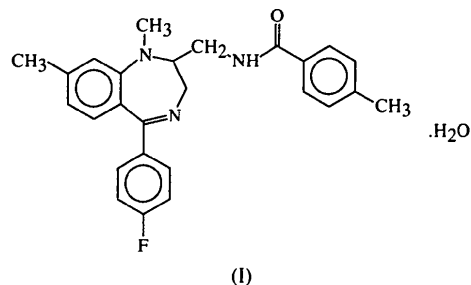
Abstract

The title compound, *N*-{[5-(4-fluorophenyl)-1,8-dimethyl-2,3-dihydro-1*H*-1,4-benzodiazepin-2-yl]methyl}-*p*-toluamide monohydrate, $C_{26}H_{26}FN_3O \cdot H_2O$, a benzodiazepine derivative with κ -opioid activity, crystallizes as a hydrate with two almost identical molecules in the asymmetric unit. The observed conformation, stabilized by two hydrogen bonds involving the H_2O molecule, is common for the 2-(acylaminoethyl)benzodiazepines. Hydrogen bonds between H_2O molecules and amidic O atoms link the non-equivalent molecules, with formation of endless chains in the *a* direction.

Comment

The title compound, (I), belongs to a series of 2-(acylaminoethyl)benzodiazepine derivatives with κ -opioid activity. Apart from the structure of tifuladom hydrate (Codding, Zeugner & Finner, 1987), the structures of this series were obtained as either chloride or toluene-sulfonate salts (Petcher, Widmer, Maetzel & Zeugner, 1985; Blaton, Peeters & De Ranter, 1996, and references therein). As both the protonation and the ionic

crystalline environment may influence the conformation of the molecule, the structure determination of a free base seems worthwhile. Unfortunately, the title compound crystallizes as a hydrate and the H_2O molecule mimics the anion site of the salts, as can be seen from the conformation given in Fig. 1.



The two molecules in the asymmetric unit of (I) are almost identical, as can be deduced from the geometric parameters of the molecules (Table 2) and the puckering parameters of the diazepine rings [$q_2 = 0.798$ (5) and 0.797 (5); $q_3 = 0.226$ (6) and 0.233 (6); $Q_T = 0.829$ (6) and 0.830 (6) Å; $\varphi_2 = -26.0$ (4) and 154.9 (3); $\varphi_3 = -128$ (1) and 51 (1); $\theta_2 = 74.2$ (4) and 73.8 (4)°, for molecules *A* and *B*, respectively, considering the atomic sequence N1—C2—C3—N4—C5—C5A—C9A]. The geometric and puckering parameters are close to those of tifuladom hydrate. The main difference between the hydrates and the salts is the value of the endocyclic angle of the N4 atom. Protonation causes the angle to open by *ca* 8°. The diazepine ring exhibits a boat conformation flattened at the stern with a pseudosymmetry plane through the C3 atom [asymmetry parameter $\Delta C_3(C3) = 0.004$ (2) and 0.008 (2) for *A* and *B*, respectively]. Each H_2O molecule is hydrogen bonded with the N4 and N12 atoms of the same

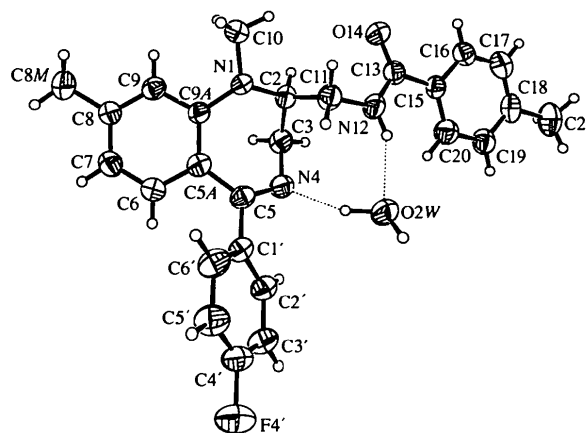


Fig. 1. Perspective view of the title compound (molecule *A*) with the atomic numbering scheme. Displacement ellipsoids are drawn at the 40% probability level.

molecule, stabilizing the observed conformation of the acylaminomethyl side chain. A third hydrogen bond between the H₂O molecule and the amidic O atom of a non-equivalent molecule forms endless chains in the a direction.

Experimental

Crystals of the title compound were obtained by slow evaporation at room temperature from a methanol/H₂O solution.

Crystal data

C₂₆H₂₆FN₃O.H₂O

M_r = 433.51

Monoclinic

*P*2₁/*n*

a = 13.6293 (6) Å

b = 10.364 (1) Å

c = 33.859 (2) Å

β = 100.236 (3)°

V = 4706.5 (5) Å³

Z = 8

D_x = 1.224 Mg m⁻³

D_m not measured

Cu Kα radiation

λ = 1.54184 Å

Cell parameters from 38

reflections

θ = 10–24°

μ = 0.680 mm⁻¹

T = 293 K

Plate

0.60 × 0.35 × 0.05 mm

Pale yellow

Data collection

Siemens P4 four-circle diffractometer

ω/2θ scans

Absorption correction:

ψ scans (*XEMP*; Siemens, 1989)

T_{min} = 0.86, *T_{max}* = 0.97

8769 measured reflections

6386 independent reflections

2130 observed reflections

[*I* > 3σ(*I*)]

R_{int} = 0.0381

θ_{max} = 57.19°

h = -1 → 14

k = -1 → 11

l = -36 → 36

3 standard reflections

monitored every 100

reflections

intensity decay: <3%

Refinement

Refinement on *F*²

R(*F*) = 0.0532

ω*R*(*F*²) = 0.1799

S = 0.853

6384 reflections

584 parameters

H-atom parameters not refined

ω = 1/[σ²(*F_o*²)]

(Δ/σ)_{max} = 0.109

Δρ_{max} = 0.30 e Å⁻³

Δρ_{min} = -0.22 e Å⁻³

Extinction correction:

SHELXL93 (Sheldrick, 1993)

Extinction coefficient:

0.00087 (9)

Atomic scattering factors

from *International Tables for X-ray Crystallography*

(1974, Vol. IV, Tables

2.2B and 2.3.1)

Table 1. Fractional atomic coordinates and equivalent isotropic displacement parameters (Å²)

$$U_{eq} = (1/3)\sum_i\sum_j U_{ij}a_i^*a_j^*a_i\cdot a_j$$

	<i>x</i>	<i>y</i>	<i>z</i>	<i>U_{eq}</i>
N1A	0.8237 (3)	0.1517 (5)	0.3396 (1)	0.057 (2)
C2A	0.8589 (4)	0.0719 (5)	0.3090 (2)	0.056 (2)
C3A	0.9139 (4)	-0.0472 (6)	0.3261 (2)	0.061 (2)

N4A	1.0132 (3)	-0.0207 (5)	0.3487 (1)	0.059 (2)
C5A	1.0160 (4)	0.0308 (5)	0.3837 (2)	0.053 (2)
C5AA	0.9283 (4)	0.0653 (5)	0.4014 (2)	0.053 (2)
C6A	0.9363 (4)	0.0472 (6)	0.4432 (2)	0.061 (2)
C7A	0.8614 (4)	0.0773 (6)	0.4633 (2)	0.065 (2)
C8A	0.7753 (4)	0.1342 (5)	0.4438 (2)	0.054 (2)
C8MA	0.6931 (5)	0.1693 (7)	0.4654 (2)	0.079 (3)
C9A	0.7646 (4)	0.1535 (5)	0.4029 (2)	0.052 (2)
C9AA	0.8394 (4)	0.1203 (5)	0.3806 (1)	0.048 (2)
C10A	0.7361 (4)	0.2291 (6)	0.3233 (2)	0.076 (3)
C11A	0.9204 (4)	0.1572 (6)	0.2858 (2)	0.061 (2)
N12A	0.9535 (3)	0.0902 (5)	0.2528 (1)	0.064 (2)
C13A	0.8925 (4)	0.0692 (6)	0.2175 (2)	0.055 (2)
O14A	0.8054 (3)	0.1058 (4)	0.2115 (1)	0.071 (2)
C15A	0.9370 (4)	-0.0016 (5)	0.1867 (1)	0.050 (2)
C16A	0.8905 (4)	0.0097 (6)	0.1472 (2)	0.061 (2)
C17A	0.9299 (5)	-0.0491 (6)	0.1168 (2)	0.069 (2)
C18A	1.0161 (5)	-0.1220 (6)	0.1255 (2)	0.065 (2)
C19A	1.0614 (4)	-0.1352 (6)	0.1650 (2)	0.066 (2)
C20A	1.0228 (4)	-0.0756 (6)	0.1954 (2)	0.065 (2)
C21A	1.0607 (5)	-0.1852 (6)	0.0922 (2)	0.090 (3)
C1'A	1.1184 (4)	0.0501 (5)	0.4078 (2)	0.054 (2)
C2'A	1.1911 (4)	-0.0420 (6)	0.4058 (2)	0.059 (2)
C3'A	1.2860 (4)	-0.0284 (6)	0.4282 (2)	0.065 (2)
C4'A	1.3064 (4)	0.0769 (6)	0.4521 (2)	0.063 (2)
F4'A	1.3994 (2)	0.0893 (4)	0.4741 (1)	0.089 (1)
C5'A	1.2381 (4)	0.1712 (6)	0.4543 (2)	0.070 (2)
C6'A	1.1425 (4)	0.1568 (6)	0.4324 (2)	0.066 (2)
N1B	1.2520 (3)	0.6483 (4)	0.3380 (1)	0.058 (2)
C2B	1.1903 (4)	0.5692 (5)	0.3071 (1)	0.053 (2)
C3B	1.1482 (4)	0.4494 (6)	0.3240 (2)	0.059 (2)
N4B	1.0687 (3)	0.4779 (4)	0.3462 (1)	0.058 (2)
C5B	1.0969 (4)	0.5274 (5)	0.3813 (2)	0.052 (2)
C5AB	1.2003 (4)	0.5611 (5)	0.3996 (1)	0.047 (2)
C6B	1.2281 (4)	0.5382 (5)	0.4408 (1)	0.056 (2)
C7B	1.3207 (4)	0.5701 (5)	0.4618 (2)	0.059 (2)
C8B	1.3888 (4)	0.6303 (5)	0.4428 (2)	0.055 (2)
C8MB	1.4893 (4)	0.6693 (7)	0.4648 (2)	0.079 (3)
C9B	1.3645 (4)	0.6508 (5)	0.4016 (1)	0.051 (2)
C9AB	1.2720 (3)	0.6174 (5)	0.3789 (1)	0.046 (2)
C10B	1.3249 (4)	0.7268 (6)	0.3219 (2)	0.074 (2)
C11B	1.1088 (4)	0.6545 (6)	0.2839 (2)	0.060 (2)
N12B	1.0473 (3)	0.5881 (4)	0.2504 (1)	0.060 (2)
C13B	1.0788 (4)	0.5640 (5)	0.2159 (2)	0.055 (2)
O14B	1.1623 (3)	0.5992 (4)	0.2104 (1)	0.069 (2)
C15B	1.0072 (4)	0.4933 (5)	0.1843 (1)	0.050 (2)
C16B	1.0219 (4)	0.5063 (6)	0.1451 (2)	0.060 (2)
C17B	0.9569 (5)	0.4478 (6)	0.1141 (2)	0.068 (2)
C18B	0.8772 (5)	0.3785 (6)	0.1214 (2)	0.068 (3)
C19B	0.8637 (4)	0.3641 (6)	0.1606 (2)	0.067 (2)
C20B	0.9283 (4)	0.4224 (6)	0.1922 (2)	0.061 (2)
C21B	0.8032 (5)	0.3155 (6)	0.0881 (2)	0.090 (3)
C1'B	1.0172 (4)	0.5478 (5)	0.4057 (2)	0.053 (2)
C2'B	0.9423 (4)	0.4589 (6)	0.4044 (2)	0.065 (2)
C3'B	0.8686 (4)	0.4740 (7)	0.4279 (2)	0.072 (2)
C4'B	0.8735 (5)	0.5808 (7)	0.4521 (2)	0.071 (3)
F4'B	0.8002 (3)	0.5933 (4)	0.4744 (1)	0.098 (2)
C5'B	0.9438 (5)	0.6710 (7)	0.4535 (2)	0.077 (3)
C6'B	1.0172 (4)	0.6551 (6)	0.4304 (2)	0.071 (2)
O1W	0.8842 (3)	0.5053 (5)	0.2876 (1)	0.092 (2)
O2W	1.1473 (3)	0.0041 (5)	0.2904 (1)	0.095 (2)

Table 2. Geometric parameters (Å, °)

N1A—C2A	1.470 (7)	N1B—C2B	1.471 (6)
N1A—C9AA	1.406 (6)	N1B—C9AB	1.399 (6)
N1A—C10A	1.462 (7)	N1B—C10B	1.464 (8)
C2A—C3A	1.506 (8)	C2B—C3B	1.522 (8)
C2A—C11A	1.530 (8)	C2B—C11B	1.523 (7)
C3A—N4A	1.458 (6)	C3B—N4B	1.456 (7)
N4A—C5A	1.294 (7)	N4B—C5B	1.287 (7)
C5A—C5AA	1.472 (8)	C5B—C5AB	1.477 (7)
C5A—C1'A	1.499 (7)	C5B—C1'B	1.492 (8)
C5AA—C6A	1.412 (7)	C5AB—C6B	1.400 (7)
C5AA—C9AA	1.408 (7)	C5AB—C9AB	1.425 (7)
C6A—C7A	1.362 (8)	C6B—C7B	1.375 (7)
C7A—C8A	1.373 (7)	C7B—C8B	1.372 (8)
C8A—C8MA	1.487 (9)	C8B—C8MB	1.494 (7)

C8A—C9A	1.382 (7)	C8B—C9B	1.391 (7)	N4A—C5A—C5AA—C9AA	-39.7 (8)
C9A—C9AA	1.415 (8)	C9B—C9AB	1.398 (6)	C5A—C5AA—C9AA—N1A	-1.0 (8)
C11A—N12A	1.453 (7)	C11B—N12B	1.457 (6)	C2A—C11A—N12A—C13A	78.5 (6)
N12A—C13A	1.347 (6)	N12B—C13B	1.339 (7)	C11A—N12A—C13A—C15A	-179.6 (4)
C13A—O14A	1.229 (7)	C13B—O14B	1.241 (7)	N12A—C13A—C15A—C16A	-160.0 (5)
C13A—C15A	1.488 (8)	C13B—C15B	1.503 (7)	C2B—N1B—C9AB—C5AB	-38.1 (7)
C15A—C16A	1.381 (7)	C15B—C16B	1.386 (7)	C10B—N1B—C9AB—C9B	-6.6 (7)
C15A—C20A	1.385 (7)	C15B—C20B	1.368 (8)	C9AB—N1B—C2B—C11B	126.0 (5)
C16A—C17A	1.386 (9)	C16B—C17B	1.385 (7)	C9AB—N1B—C2B—C3B	0.8 (7)
C17A—C18A	1.383 (9)	C17B—C18B	1.361 (9)	N1B—C2B—C11B—N12B	175.8 (4)
C18A—C19A	1.376 (8)	C18B—C19B	1.382 (9)	N1B—C2B—C3B—N4B	72.7 (6)
C18A—C21A	1.521 (9)	C18B—C21B	1.521 (8)	C2B—C3B—N4B—C5B	-74.8 (6)
C19A—C20A	1.384 (9)	C19B—C20B	1.396 (8)	C3B—N4B—C5B—C5AB	2.4 (8)
C1'A—C2'A	1.386 (8)	C1'B—C2'B	1.370 (8)	N4B—C5B—C1'B—C2'B	37.8 (7)
C1'A—C6'A	1.388 (8)	C1'B—C6'B	1.393 (8)	N4B—C5B—C5AB—C9AB	39.4 (8)
C2'A—C3'A	1.385 (7)	C2'B—C3'B	1.397 (9)	C5B—C5AB—C9AB—N1B	0.3 (8)
C3'A—C4'A	1.356 (9)	C3'B—C4'B	1.370 (9)	C2B—C11B—N12B—C13B	-76.1 (6)
C4'A—F4'A	1.358 (6)	C4'B—F4'B	1.363 (8)	N12B—C13B—C15B—C16B	158.3 (5)
C4'A—C5'A	1.362 (9)	C4'B—C5'B	1.334 (9)		
C5'A—C6'A	1.388 (8)	C5'B—C6'B	1.383 (9)		

C9AA—N1A—C10A	118.0 (4)	C9AB—N1B—C10B	118.1 (4)
C2A—N1A—C10A	113.0 (4)	C2B—N1B—C10B	112.6 (4)
C2A—N1A—C9AA	123.7 (4)	C2B—N1B—C9AB	124.3 (4)
N1A—C2A—C11A	108.3 (4)	N1B—C2B—C11B	108.4 (4)
N1A—C2A—C3A	113.1 (4)	N1B—C2B—C3B	113.1 (4)
C3A—C2A—C11A	113.0 (4)	C3B—C2B—C11B	112.3 (4)
C2A—C3A—N4A	113.6 (5)	C2B—C3B—N4B	113.2 (4)
C3A—N4A—C5A	115.5 (4)	C3B—N4B—C5B	115.4 (4)
N4A—C5A—C1'A	115.2 (4)	N4B—C5B—C1'B	116.0 (4)
N4A—C5A—C5AA	125.3 (5)	N4B—C5B—C5AB	126.1 (5)
C5AA—C5A—C1'A	119.4 (4)	C5AB—C5B—C1'B	117.9 (4)
C5A—C5AA—C9AA	125.8 (4)	C5B—C5AB—C9AB	125.2 (4)
C5A—C5AA—C6A	116.8 (4)	C5B—C5AB—C6B	116.6 (4)
C6A—C5AA—C9AA	117.4 (5)	C6B—C5AB—C9AB	118.2 (4)
C5AA—C6A—C7A	122.8 (5)	C5AB—C6B—C7B	122.4 (5)
C6A—C7A—C8A	120.6 (5)	C6B—C7B—C8B	120.1 (5)
C7A—C8A—C9A	118.3 (5)	C7B—C8B—C9B	118.7 (5)
C7A—C8A—C8MA	121.6 (5)	C7B—C8B—C8MB	122.6 (5)
C8MA—C8A—C9A	120.1 (5)	C8MB—C8B—C9B	119.6 (5)
C8A—C9A—C9AA	122.8 (5)	C8B—C9B—C9AB	123.0 (4)
C5AA—C9AA—C9A	118.1 (4)	C5AB—C9AB—C9B	117.4 (4)
N1A—C9AA—C9A	118.3 (4)	N1B—C9AB—C9B	119.0 (4)
N1A—C9AA—C5AA	123.5 (4)	N1B—C9AB—C5AB	123.4 (4)
C2A—C11A—N12A	113.1 (4)	C2B—C11B—N12B	113.1 (4)
C11A—N12A—C13A	122.2 (4)	C11B—N12B—C13B	122.5 (4)
N12A—C13A—C15A	115.9 (4)	N12B—C13B—C15B	116.4 (4)
N12A—C13A—O14A	121.5 (5)	N12B—C13B—O14B	121.5 (5)
O14A—C13A—C15A	122.6 (5)	O14B—C13B—C15B	122.2 (5)
C13A—C15A—C20A	123.9 (4)	C13B—C15B—C20B	123.8 (4)
C13A—C15A—C16A	117.5 (5)	C13B—C15B—C16B	116.7 (5)
C16A—C15A—C20A	118.6 (5)	C16B—C15B—C20B	119.5 (5)
C15A—C16A—C17A	120.7 (5)	C15B—C16B—C17B	120.0 (5)
C16A—C17A—C18A	120.6 (5)	C16B—C17B—C18B	121.2 (5)
C17A—C18A—C21A	120.9 (5)	C17B—C18B—C21B	122.5 (5)
C17A—C18A—C19A	118.6 (6)	C17B—C18B—C19B	118.6 (6)
C19A—C18A—C21A	120.5 (6)	C19B—C18B—C21B	118.9 (5)
C18A—C19A—C20A	121.0 (6)	C18B—C19B—C20B	121.1 (5)
C15A—C20A—C19A	120.5 (5)	C15B—C20B—C19B	119.5 (5)
C5A—C1'A—C6'A	122.1 (5)	C5B—C1'B—C6'B	121.6 (5)
C5A—C1'A—C2'A	119.1 (5)	C5B—C1'B—C2'B	119.9 (5)
C2'A—C1'A—C6'A	118.8 (5)	C2'B—C1'B—C6'B	118.4 (5)
C1'A—C2'A—C3'A	120.7 (5)	C1'B—C2'B—C3'B	120.9 (6)
C2'A—C3'A—C4'A	118.7 (5)	C2'B—C3'B—C4'B	117.8 (6)
C3'A—C4'A—C5'A	122.7 (6)	C3'B—C4'B—C5'B	123.1 (6)
C3'A—C4'A—F4'A	118.5 (5)	C3'B—C4'B—F4'B	116.5 (6)
F4'A—C4'A—C5'A	118.8 (5)	F4'B—C4'B—C5'B	120.4 (6)
C4'A—C5'A—C6'A	118.7 (6)	C4'B—C5'B—C6'B	118.9 (6)
C1'A—C6'A—C5'A	120.4 (5)	C1'B—C6'B—C5'B	120.8 (6)

C2A—N1A—C9AA—C5AA	38.6 (7)
C10A—N1A—C9AA—C9A	6.5 (7)
C9AA—N1A—C2A—C11A	-126.2 (5)
C9AA—N1A—C2A—C3A	-0.1 (7)
N1A—C2A—C11A—N12A	-176.1 (4)
N1A—C2A—C3A—N4A	-73.2 (6)
C2A—C3A—N4A—C5A	74.1 (6)
C3A—N4A—C5A—C5AA	-1.0 (8)
N4A—C5A—C1'A—C2'A	-37.1 (7)

Table 3. Hydrogen-bonding geometry (Å, °)

D—H...A	D—H	H...A	D...A	D—H...A
N12A—H12A...O2W	0.860	2.07	2.863 (5)	153
O2W—H2WB...N4A	1.061	1.92	2.930 (6)	159
O2W—H2WA...O14B'	1.079	1.76	2.781 (5)	155
N12B—H12B...O1W	0.860	2.08	2.873 (6)	152
O1W—H1WB...N4B	1.199	1.97	2.929 (5)	133
O1W—H1WA...O14A''	1.170	1.74	2.791 (5)	147

Symmetry codes: (i) $\frac{1}{2} - x, y - \frac{1}{2}, \frac{1}{2} - z$; (ii) $\frac{3}{2} - x, \frac{1}{2} + y, \frac{1}{2} - z$.

The title structure was solved by direct methods and refined by full-matrix least squares on F^2 . H atoms (except those of the H₂O molecules obtained from the ΔF synthesis) were included at calculated positions riding on their parent atoms.

Data collection: XSCANS (Siemens, 1994). Cell refinement: XSCANS. Data reduction: XSCANS. Program(s) used to solve structure: SIR92 (Altomare *et al.*, 1994). Program(s) used to refine structure: SHELXL93 (Sheldrick, 1993). Molecular graphics: ORTEX (McArdle, 1994). Software used to prepare material for publication: PARST (Nardelli, 1983).

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Lists of structure factors, anisotropic displacement parameters, H-atom coordinates and complete geometry have been deposited with the IUCR (Reference: NA1261). Copies may be obtained through The Managing Editor, International Union of Crystallography, 5 Abbey Square, Chester CH1 2HU, England.

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