

C(7)—O—C(8)	101.8 (2)	C(1)—C(6)—C(5)	119.2 (2)
C(4)—N(1)—C(7)	125.7 (3)	O—C(7)—N(1)	115.7 (2)
N(3)—N(2)—C(7)	105.0 (2)	O—C(7)—N(2)	113.6 (3)
N(2)—N(3)—C(8)	107.9 (2)	N(1)—C(7)—N(2)	130.8 (2)
C(11)—N(4)—C(12)	116.3 (3)	O—C(8)—N(3)	111.8 (3)
Cl—C(1)—C(2)	119.7 (2)	O—C(8)—C(9)	120.6 (2)
Cl—C(1)—C(6)	119.3 (2)	N(3)—C(8)—C(9)	127.6 (2)
C(2)—C(1)—C(6)	121.0 (2)	C(8)—C(9)—C(10)	123.6 (2)
C(1)—C(2)—C(3)	119.6 (2)	C(8)—C(9)—C(13)	118.7 (3)
C(2)—C(3)—C(4)	120.5 (2)	C(10)—C(9)—C(13)	117.8 (2)
N(1)—C(4)—C(3)	116.9 (3)	C(9)—C(10)—C(11)	119.6 (2)
N(1)—C(4)—C(5)	124.3 (2)	N(4)—C(11)—C(10)	123.6 (2)
C(3)—C(4)—C(5)	118.9 (2)	N(4)—C(12)—C(13)	124.0 (2)
C(4)—C(5)—C(6)	120.8 (2)	C(9)—C(13)—C(12)	118.7 (2)

Data collection was performed using CAD-4 diffractometer software (Enraf–Nonius, 1977). The structure was solved by direct methods with the MULTAN11/82 programs (Main *et al.*, 1982). The structure was refined by full-matrix least-squares methods using the SDP programs (Frenz, 1978).

Lists of structure factors, anisotropic displacement parameters, H-atom coordinates, bond distances and angles involving H atoms, least-squares-planes data and torsion angles have been deposited with the IUCr (Reference: AB1133). Copies may be obtained through The Managing Editor, International Union of Crystallography, 5 Abbey Square, Chester CHI 2HU, England.

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Depogen, a Haemorheological Agent

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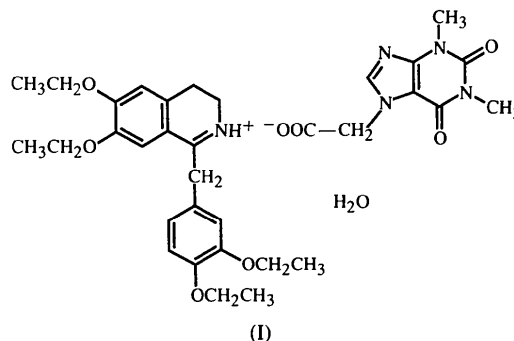
(Received 28 November 1994; accepted 13 January 1995)

Abstract

The crystal structure of the haemorheological agent depogen {1-[(3,4-diethoxyphenyl)methyl]-6,7-diethoxy-3,4-dihydro-2H⁺-isoquinolinium 1,2,3,6-tetrahydro-7H-1,3-dimethyl-2,6-dioxopurine-7-acetate monohydrate, C₂₄H₃₂NO₄⁺.C₉H₉N₄O₄⁻.H₂O} reveals the role of a water molecule in building up a network of hydrogen bonds in its crystal lattice. Knowledge of the structure may help the elucidation of the differences in the pharmacological properties of the hydrated and the dehydrated compound.

Comment

The haemorheological agent depogen, (I) (Kapui *et al.*, 1992), has been produced in two crystal forms. While the cotton-like anhydrous form resisted attempts to produce single crystals of good quality (and indeed a substance suitable for drug formulation purposes), the monohydrate crystals were of sufficient quality to be subjected to X-ray diffractometry.



The N-atom-containing ring of the isoquinoline moiety in the cation is in a screw boat conformation (2S_3), as shown by the sign combination of the corresponding torsion angles (0, +, -, +, 0, -) (Table 2). Apart from the ethoxy group containing the disordered C29B atom, the remaining three are closely embedded in the plane of their corresponding benzene ring.

It is worth mentioning that the shape of the cation is characterized by the torsion angles N1—C2—C11—C12 and C2—C11—C12—C13. Accordingly, in other crystal structures containing the same cation but only a single chloride anion and different solvent molecules, these two torsion angles vary the most, while other structural parameters (such as the freely rotating torsion angles involving the atoms of the ethoxy groups) remain essentially the same in all the examined structures (Böcskei, Menyhárd, Simon, Friesz & Hermeicz, 1994).

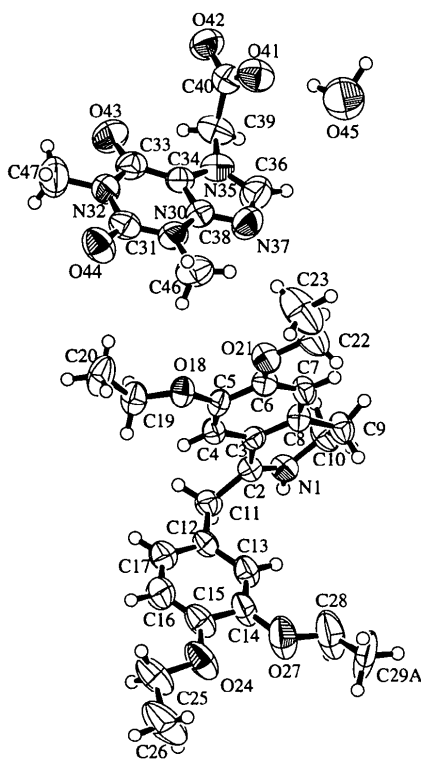


Fig. 1. Molecular structure and atomic numbering of the title compound.

As far as the intermolecular contacts are concerned, it can be seen (Fig. 2) that the water molecule plays an essential role in the construction of the crystal structure, since it bridges the O41 and O42 atoms of the carboxylate groups of two separate theophylline-7-acetate residues. Additionally, O41 is the acceptor atom in another hydrogen bond donated by N1 of the isoquinoline residue. The graph notation (Etter, 1990) for the hydrogen-bond chain consisting of the hydrogen bonds donated by the water molecules is $C^22(6)$, while the symbol for that donated by the N1 atom is D .

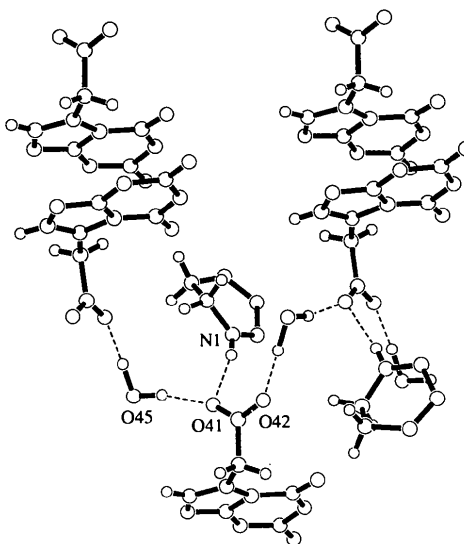


Fig. 2. Hydrogen-bonding interactions in the crystal structure of the title compound.

The angle formed by the two hydrogen bonds connected to O41 is *ca* 90° , somewhat narrower than expected from the possible localization of the non-bonding electron pairs.

Experimental

Crystal data

$C_{24}H_{32}NO_4^+ \cdot C_9H_9N_4O_4^- \cdot H_2O$

$M_r = 653.72$

Orthorhombic

$Fdd2$

$a = 24.612(6) \text{ \AA}$

$b = 60.310(8) \text{ \AA}$

$c = 9.131(7) \text{ \AA}$

$V = 13\,553(7) \text{ \AA}^3$

$Z = 16$

$D_x = 1.281 \text{ Mg m}^{-3}$

Cu $K\alpha$ radiation

$\lambda = 1.5418 \text{ \AA}$

Cell parameters from 20 reflections

$\theta = 20.93\text{--}57.85^\circ$

$\mu = 0.779 \text{ mm}^{-1}$

$T = 293(2) \text{ K}$

Hexagonal

$0.300 \times 0.300 \times 0.150 \text{ mm}$

Yellow

Data collection

AFC-6S diffractometer

$\omega/2\theta$ scans

Absorption correction:
none

4311 measured reflections

3738 independent reflections

1925 observed reflections

$[I > 2\sigma(I)]$

$\theta_{\max} = 75.36^\circ$

$h = -30 \rightarrow 29$

$k = -54 \rightarrow 75$

$l = -6 \rightarrow 11$

3 standard reflections

monitored every 150

reflections

intensity decay: 0.95%

Refinement

Refinement on F^2

$R(F) = 0.0581$

$wR(F^2) = 0.1556$

Extinction correction:

SHELXL93 (Sheldrick,

1993)

$S = 1.126$
 3733 reflections
 439 parameters
 Only H-atom U 's refined
 $w = 1/[\sigma^2(F_o^2) + (0.1162P)^2 + 6.9099P]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} = -0.240$
 $\Delta\rho_{\max} = 0.341 \text{ e } \text{Å}^{-3}$
 $\Delta\rho_{\min} = -0.297 \text{ e } \text{Å}^{-3}$

Extinction coefficient:
 0.00006 (2)
 Atomic scattering factors
 from *International Tables*
 for *Crystallography* (1992),
 Vol. C, Tables 4.2.6.8 and
 6.1.1.4)
 Absolute configuration:
 Flack (1983)

Table 1. *Fractional atomic coordinates and equivalent isotropic displacement parameters* (Å^2)

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

	x	y	z	U_{eq}
N1	0.7246 (2)	0.02130 (10)	-0.1008 (8)	0.056 (2)
C2	0.7296 (3)	0.03492 (11)	0.0082 (9)	0.048 (2)
C3	0.6872 (3)	0.05122 (11)	0.0354 (8)	0.046 (2)
C4	0.6858 (3)	0.06326 (11)	0.1662 (8)	0.049 (2)
C5	0.6470 (3)	0.07943 (11)	0.1854 (8)	0.047 (2)
C6	0.6103 (3)	0.08436 (11)	0.0710 (9)	0.051 (2)
C7	0.6122 (3)	0.07243 (12)	-0.0581 (9)	0.056 (2)
C8	0.6502 (3)	0.05577 (11)	-0.0762 (9)	0.051 (2)
C9	0.6542 (3)	0.04326 (13)	-0.2175 (10)	0.063 (2)
C10	0.6763 (4)	0.02038 (14)	-0.1911 (11)	0.069 (2)
C11	0.7801 (3)	0.03327 (12)	0.0994 (10)	0.058 (2)
C12	0.8161 (3)	0.05343 (12)	0.0891 (9)	0.053 (2)
C13	0.8310 (3)	0.06161 (13)	-0.0471 (10)	0.066 (2)
C14	0.8649 (4)	0.07943 (14)	-0.0593 (12)	0.072 (3)
C15	0.8862 (4)	0.08926 (14)	0.0668 (12)	0.074 (3)
C16	0.8717 (4)	0.0812 (2)	0.1997 (12)	0.084 (3)
C17	0.8374 (4)	0.0631 (2)	0.2105 (12)	0.079 (3)
O18	0.6406 (2)	0.09182 (8)	0.3094 (6)	0.0575 (13)
C19	0.6651 (4)	0.08420 (13)	0.4400 (10)	0.066 (2)
C20	0.6437 (5)	0.0981 (2)	0.5642 (11)	0.093 (3)
O21	0.5753 (2)	0.10120 (9)	0.0993 (7)	0.0644 (14)
C22	0.5366 (4)	0.1071 (2)	-0.0117 (14)	0.092 (4)
C23	0.5034 (5)	0.1253 (2)	0.052 (2)	0.134 (6)
O24	0.9210 (3)	0.10661 (10)	0.0435 (9)	0.097 (2)
C25	0.9481 (4)	0.1154 (2)	0.1732 (13)	0.099 (4)
C26	0.9895 (5)	0.1322 (2)	0.1178 (17)	0.148 (7)
O27	0.8807 (4)	0.08854 (15)	-0.1880 (9)	0.123 (3)
C28	0.8674 (5)	0.0775 (2)	-0.3312 (11)	0.146 (7)
C29A	0.8724 (5)	0.0959 (2)	-0.4360 (11)	0.16 (2)
C29B	0.8150 (8)	0.0893 (4)	-0.3717 (25)	0.128 (12)
N30	0.4906 (3)	0.05157 (11)	0.4346 (10)	0.068 (2)
C31	0.4935 (3)	0.05137 (14)	0.5840 (12)	0.069 (2)
N32	0.4673 (3)	0.03396 (13)	0.6564 (9)	0.071 (2)
C33	0.4425 (3)	0.0155 (2)	0.5877 (12)	0.069 (2)
C34	0.4429 (3)	0.01724 (13)	0.4361 (10)	0.057 (2)
N35	0.4248 (3)	0.00283 (11)	0.3272 (10)	0.071 (2)
C36	0.4370 (5)	0.0128 (2)	0.2008 (14)	0.093 (3)
N37	0.4635 (4)	0.0326 (2)	0.2153 (10)	0.092 (3)
C38	0.4663 (3)	0.03480 (14)	0.3622 (10)	0.064 (2)
C39	0.3935 (3)	-0.01708 (13)	0.3485 (14)	0.078 (3)
C40	0.3329 (3)	-0.01352 (14)	0.3605 (11)	0.062 (2)
O41	0.3141 (2)	0.00163 (11)	0.2897 (10)	0.090 (2)
O42	0.3071 (3)	-0.02675 (12)	0.4371 (9)	0.090 (2)
O43	0.4232 (3)	0.00046 (12)	0.6625 (9)	0.100 (2)
O44	0.5167 (3)	0.06593 (11)	0.6534 (9)	0.092 (2)
O45	0.3015 (5)	0.0307 (2)	0.0355 (13)	0.165 (5)
C46	0.5146 (4)	0.0703 (2)	0.3559 (15)	0.101 (4)
C47	0.4695 (6)	0.0338 (2)	0.8164 (13)	0.110 (4)

Table 2. *Selected bond lengths* (Å) *and torsion angles* ($^\circ$)

N1—C2	1.296 (9)	N1—C10	1.447 (10)
C10—N1—C2—C3	-4.9 (11)	C8—C9—C10—N1	-48.6 (10)
N1—C2—C3—C8	-16.9 (10)	N1—C2—C11—C12	114.3 (7)
C4—C3—C8—C9	177.4 (7)	C2—C11—C12—C13	-51.0 (10)
C3—C8—C9—C10	30.6 (10)	C34—N35—C39—C40	84.1 (12)
C2—N1—C10—C9	38.6 (11)	N35—C39—C40—O41	32.7 (14)

Table 3. *Hydrogen-bonding geometry* (Å , $^\circ$)

$D-H \cdots A$	$D \cdots A$	$D-H \cdots A$
O45—H45A \cdots O41	2.924 (13)	168.1 (4)
O45'—H45B' \cdots O42	2.828 (13)	174.9 (5)
N1''—H1 \cdots O41	2.695 (9)	158.9 (3)

Table 4. *Least-squares-planes data*

Equation of plane: $15.600 (35)x + 40.873 (65)y - 3.404 (16)z = 12.720 (22)$
 Deviation of atoms from the plane (Å):
 (* indicates atom used to define plane)

C2*	0.061 (5)	O18*	-0.027 (5)
C3*	-0.026 (6)	O21*	0.053 (5)
C4*	-0.001 (6)	C19	-0.401 (10)
C5*	-0.011 (6)	C20	-0.590 (13)
C6*	0.007 (6)	C22	0.068 (15)
C7*	-0.011 (7)	C23	0.076 (19)
C8*	-0.038 (7)	N1	-0.203 (8)
C9*	-0.006 (6)	C10	-0.685 (12)

R.m.s. deviation of fitted atoms: 0.031 Å

Equation of plane: $19.316 (30)x - 37.331 (84)y - 0.277 (34)z = 13.767 (31)$

Angle to previous plane (with approximate e.s.d.): $84.88 (13)^\circ$

Deviation of atoms from the plane (Å):
 (* indicates atom used to define plane)

C11*	0.033 (5)	O24*	0.032 (6)
C12*	-0.022 (6)	O27*	-0.008 (7)
C13*	-0.001 (8)	C25	0.190 (15)
C14*	-0.009 (9)	C26	0.379 (19)
C15*	0.000 (9)	C28	0.189 (19)
C16*	-0.017 (8)	C29A	-0.375 (21)
C17*	-0.007 (8)	C29B	-1.253 (23)

R.m.s. deviation of fitted atoms: 0.018 Å

Equation of plane: $21.396 (23)x - 29.779 (77)y - 0.200 (18)z = 8.893 (13)$

Angle to previous plane (with approximate e.s.d.): $8.7 (2)^\circ$

Deviation of atoms from the plane (Å):
 (* indicates atom used to define plane)

N30*	-0.018 (7)	C38*	-0.025 (8)
C31*	0.020 (8)	C39*	-0.034 (6)
N32*	-0.036 (8)	O43*	0.017 (8)
C33*	-0.005 (8)	O44*	0.069 (7)
C34*	-0.018 (8)	C46*	-0.047 (9)
N35*	0.047 (7)	C47*	-0.016 (10)
C36*	0.035 (10)	C40	-1.439 (10)
N37*	0.011 (9)		

R.m.s. deviation of fitted atoms: 0.033 Å

Data collection: *MSC/AFC Diffractometer Control Software* (Molecular Structure Corporation, 1988). Cell refinement: *MSC/AFC Diffractometer Control Software*. Data reduction: *TEXSAN* (Molecular Structure Corporation, 1985). Program(s) used to solve structure: *SHELXS86* (Sheldrick, 1985). Program(s) used to refine structure: *SHELXL93* (Sheldrick, 1993). Software used to prepare material for publication: *TEXSAN*.

Lists of structure factors, anisotropic displacement parameters, H-atom coordinates and complete geometry have been deposited with the IUCr (Reference: KA1104). Copies may be obtained through The Managing Editor, International Union of Crystallography, 5 Abbey Square, Chester CH1 2HU, England.

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2-(2-Methoxyphenyl)pyridine 1-Oxide and its 3-Methyl Derivative

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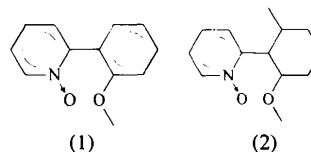
Abstract

The crystal structures of 2-(2-methoxyphenyl)pyridine 1-oxide, C₁₂H₁₁NO₂ (1), and 2-(2-methoxyphenyl)-3-methylpyridine 1-oxide, C₁₃H₁₃NO₂ (2), have been determined. The dihedral angle between the least-squares planes of the aromatic rings is 66.6 (1)° for (1) and 74.70 (6)° for (2). Weak C—H···O hydrogen bonds govern the crystal packing.

Comment

Orellanine (3,3',4,4'-tetrahydroxy-2,2'-bipyridine 1,1'-dioxide) was shown to lose both *N*-oxide groups at a surprisingly low temperature (Antkowiak & Gessner, 1979) compared with other bipyridine *N,N'*-dioxides (Wenkert & Woodward, 1983). As part of our studies on the mechanism of this deoxygenation reaction, we reported the structure of a model compound, 2-(2-hydroxyphenyl)pyridine 1-oxide (Kubicki, Borowiak, Antkowiak & Antkowiak, 1990), which has all the structural features necessary for the proposed mechanism of this deoxygenation process (Antkowiak & Gessner, 1984). In this structure, we found a strong intramolecular O—H···O—N hydrogen bond and the dihedral angle between the two aromatic rings was 38.2 (1)°. Since decrease of the twist angle is supposed to play a crucial role in the transition state of the deoxygenation process, we decided to check the influence of intramolecular hydrogen bonding on the molecular conformation by studying related compounds having no possibility of

forming such a bond. In this paper, we present the structure determinations of two compounds, 2-(2-methoxyphenyl)pyridine 1-oxide, (1), and its 3-methyl derivative, (2).



The twist angle between the two almost perfectly planar aromatic rings is 66.6 (1)° in (1) and 74.70 (6)° in (2). This seems to prove the role of intramolecular hydrogen bonding in stabilizing the conformation, reflected in a smaller dihedral angle, for this class of compounds. Comparing the conformations of (1) and (2), it was found that the rings were twisted in opposite directions; the absolute values of the torsion angles N1—C2—C1'—C2' are 69.2 (3)° and 106.8 (2)° in compounds (1) and (2), respectively. Thus, if one imagines a plane perpendicular to the pyridine ring and containing the central C2—C1' bond, in (1) both O atoms will be on the same side of this plane, while in (2) they will lie on opposite sides. In the case of (1), even despite the lack of steric hindrance at positions 3 and 6', a relatively short non-bonded O1···O2' distance of 3.032 (3) Å is found. A similar situation is observed in 2,2'-bipyridine 1,1'-dioxide, where the absolute value of the torsion angle N1—C2—C2'—N1' is 70.9 (2)° and the O1···O1' distance is 3.040 (2) Å (Dutkiewicz, 1995). It is also noteworthy that the crystal structure of (2) contains both conformational enantiomers, while in (1) there are only homochiral molecules. The disposition of the methoxy group with respect to the phenyl ring is also slightly different in both compounds. In (1), the methoxy group is almost coplanar with the phenyl ring, while in (2) it is significantly tilted out of the plane; the torsion angles C3'—C2'—O2'—C2'1 are -3.3 (5)° and -15.9 (3)° for compounds (1) and (2), respectively. This is accompanied by a statistically significant difference in the values of the C—O—C bond angles: 118.1 (3)° for (1) and 116.5 (2)° for (2). The introduction of the methyl substituent in the 3-position changes the bond-angle pattern in the pyridine ring.

The differences between (2) and (1) are in agreement with the observations of Domenicano & Murray-Rust (1979) on mono- and disubstituted benzene derivatives. The N—O bond lengths of 1.307 (4) Å for (1) and 1.309 (2) Å for (2) are close to the typical values for N—O groups not involved in hydrogen bonding (Eichhorn, 1987).

In both structures, the molecular packing is determined by C—H···O hydrogen bonds. Compound (1) has a two-dimensional structure with chains of molecules along the [100] and [010] directions, interconnected with the rings of four other molecules. In (2),