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A Novel Intercalation of a Water Molecule Between Pyrimidine Bases in 5-Nitro-1-(β -D-ribosyluronic acid)–Uracil Monohydrate [1-(5-Nitro-2,4-dioxypyrimidinyl)- β -D-ribofuranuronic Acid Monohydrate]*

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Abstract. $C_9H_9N_3O_9 \cdot H_2O$, monoclinic, $P2_1$, $a = 8.982$ (1), $b = 10.245$ (1), $c = 6.651$ (1) Å, $\beta = 92.27$ (1)° at $22 \pm 3^\circ\text{C}$, $V = 611.5$ Å³, $Z = 2$, FW 321, $\lambda(\text{Cu } K\alpha) = 1.54051$ Å, D_m (flotation) = 1.74, $D_x = 1.75$ g cm⁻³, $\mu(\text{Cu } K\alpha) = 7.6$ cm⁻¹, $R = 0.056$. The molecule exhibits *anti* conformation ($\chi_{\text{CN}} = 53.9^\circ$), an intramolecular C(6)–H...O(5') hydrogen bond and C(2')-endo, C(3')-exo (2T_3) pucker. Two unusual features in this structure are (a) the intercalation of water molecules between stacked pyrimidine rings 6.55 Å apart and related by the *c* translation and (b) a number of C–H...O interactions.

Introduction. In connection with a project on radiation sensitization, we have been investigating a series of nitro-substituted bases and nucleosides. Our interest in these structures was to study the hydrogen-bonding scheme of these modified nucleosides as well as to correlate the orientation of the 'g' tensor of these nitro radicals (analysis in progress by Dr H. Box of this Institute) using spin resonance techniques.

The title compound (I) was first synthesized by Fox and co-workers (Wempen, Doerr, Kaplan & Fox, 1960) in the course of their screening of nitro analogs of nucleosides for possible antitumour activities. Crystals of the title compound were obtained by the addition of conc. HNO_3 to the aqueous solution of uridine, followed by slow evaporation of the resulting solution. The unit-cell and other crystallographic data are given

in the *Abstract*. 1438 reflections (only nine had intensities $<2\sigma$) to the limit of $2\theta = 165^\circ$ for Cu $K\alpha$ were collected using the stationary-crystal–stationary-counter procedure (Furnas & Harker, 1955) and were processed in the usual way. A detailed absorption correction depending on the shape of the crystal (Coppens, 1970) was carried out.

The structure (Fig. 1) was solved by a combination of *MULTAN* (Germain, Main & Woolfson, 1971) and trial-and-error methods. One of the *E* maps gave two

Table 1. Atomic coordinates ($\times 10^4$) for the non-hydrogen atoms with e.s.d.'s in parentheses

	<i>x</i>	<i>y</i>	<i>z</i>
O(2)	–9044 (3)	–1340	–2062 (5)
O(4)	–4194 (4)	–314 (4)	–2930 (7)
O(5a)	–3341 (4)	–4313 (4)	–3432 (10)
O(5b)	–2341 (4)	–2420 (4)	–2980 (7)
O(2')	–9866 (4)	–4206 (4)	–5669 (6)
O(3')	–11014 (4)	–6051 (4)	–2972 (7)
O(1')	–8172 (4)	–4886 (3)	–874 (5)
O(5')	–5929 (4)	–6404 (4)	–2014 (7)
O(5'')	–7240 (4)	–8221 (4)	–1602 (8)
OW	–6568 (4)	–2662 (4)	–7914 (7)
N(1)	–7450 (4)	–3033 (3)	–2631 (6)
N(3)	–6599 (4)	–886 (4)	–2476 (7)
N(C5)	–3429 (4)	–3132 (5)	–3147 (7)
C(2)	–7795 (5)	–1717 (4)	–2357 (7)
C(4)	–5118 (5)	–1182 (5)	–2781 (7)
C(5)	–4888 (4)	–2585 (5)	–2940 (7)
C(6)	–6043 (5)	–3426 (5)	–2863 (7)
C(1')	–8635 (4)	–4028 (4)	–2458 (7)
C(2')	–8891 (5)	–4868 (4)	–4302 (7)
C(3')	–9479 (4)	–6124 (4)	–3402 (7)
C(4')	–8545 (5)	–6215 (4)	–1449 (7)
C(5')	–7151 (5)	–7047 (4)	–1715 (7)

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Fig. 2. The molecule in the *anti* conformation. Note the intramolecular C—H...O and the O(5')—H(O5')...OW hydrogen bonding in the structure. The torsion angles around the ribose ring, and the C(6)—N(1)—C(1')—O(1') and C(6)—C(5)—N(C5)—O(5a) torsion angles are shown.

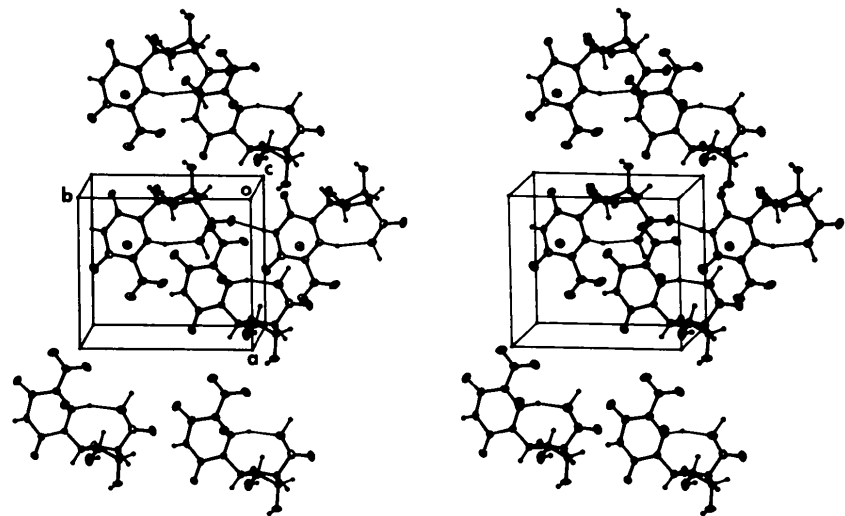


Fig. 3. A stereoview of the packing and hydrogen bonding of the molecules in the crystal structure.

Table 3. *Hydrogen bonds and weak hydrogen interactions*

<i>D</i> —H... <i>A</i>	<i>D</i> —H (Å)	H... <i>A</i> (Å)	<i>D</i> ... <i>A</i> (Å)	∠ <i>D</i> —H... <i>A</i> (°)
N(3)—H(N3)...O(5'') ⁱ	0.74 (5)	2.13 (5)	2.855 (6)	168 (5)
O(2')—H(O2')...O(2) ^{iv}	0.85 (6)	2.13 (7)	2.810 (5)	137 (6)
O(3')—H(O3')...O(5a) ^v	0.53 (6)	2.30 (6)	2.754 (5)	146 (8)
O(5')—H(O5')...O <i>W</i> ^{vi}	1.2 (1)	1.7 (1)	2.589 (5)	128 (9)
C(6)—H(C6)...O(5')	1.11 (9)	2.11 (7)	3.104 (6)	147 (6)
C(2')—H(C2')...O(4) ⁱⁱ	1.04 (5)	2.41 (5)	3.418 (6)	163 (4)
C(3')—H(C3')...O(2') ^{iv}	0.96 (7)	2.51 (7)	3.268 (6)	136 (5)
C(4')—H(C4')...O(2) ⁱⁱⁱ	1.29 (11)	2.27 (11)	3.251 (6)	130 (7)

The superscripts refer to the following symmetry relationships:
(i) $x, 1 + y, z$; (ii) $\bar{1} - x, -\frac{1}{2} + y, \bar{1} - z$; (iii) $\bar{2} - x, -\frac{1}{2} + y, \bar{z}$; (iv) $\bar{2} - x, -\frac{1}{2} + y, \bar{1} - z$; (v) $\bar{1} + x, y, z$;
(vi) $\bar{1} - x, -\frac{1}{2} + y, \bar{1} - z$.

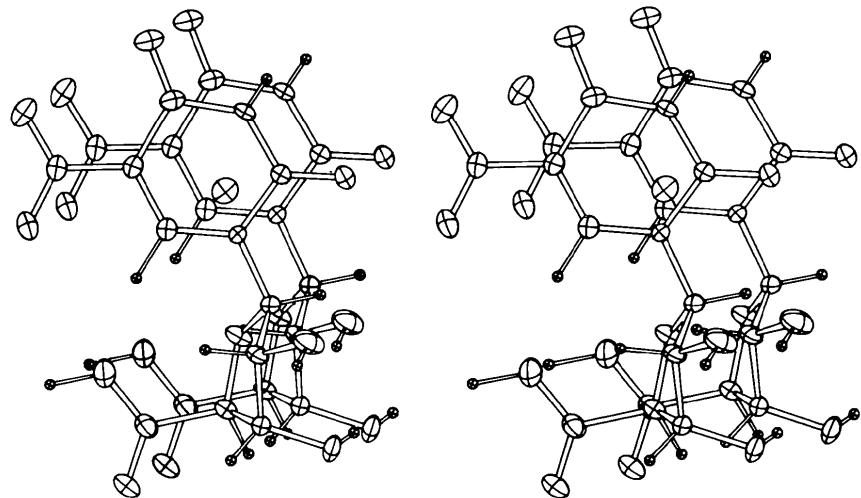


Fig. 4. A stereoview of the intercalation of the water molecule between the nucleic acid bases, viewed normal to the plane of the base.

hydrogen-bonding distances of 2.86 and 2.81 Å respectively from O(4)[$\bar{1} - x, -\frac{1}{2} + y, \bar{1} - z$] and O(3')[$2 - x, y + \frac{1}{2}, 1 - z$] and is possibly hydrogen bonded to them [O(4)···O $\bar{W}$ ···O(3'), 134°], but the location of the H atoms of the water is not unequivocal. Around the water O atom, two diffuse clouds of electron density (of approximately 0.3 e Å⁻³ extending up to 1 Å in the **b** direction) are present. If these densities are taken to represent the two H atoms of the water molecule and their coordinates are refined, their new orientations do not correspond to the possible hydrogen bonding involving O(3') and O(4).

The intercalation of water between nucleic acid bases (Fig. 4) is very unusual (Srikrishnan & Parthasarathy, 1976). It is interesting to note that 5-nitrouridine monohydrate (Egert, Lindner, Hillen & Gassen, 1977) does not form a water sandwich. The water molecules are found to be nearly in the plane of nucleic acid bases (Fig. 2), as found in other related structures. In addition, they intercalate between bases in one stack of nucleic acid bases but are in the same plane as the nucleic acid bases in an adjacent stack, that is shifted, vertically relative to the first stack, by approximately 3.2 Å.

The occurrence of a C(sp²)-H···O hydrogen bond (Table 3) [C(6)-HC(6)···O(5')] in this structure is noteworthy. The presence of an 'electron-withdrawing' nitro group adjacent to C(6)-H causes additional polarization of this bond and enables this H to take part in a hydrogen bond. In addition, there are three C(sp³)-H···O interactions (Table 3) involving the ribose C atoms. These interactions are unexpected since these contacts involve C(sp³)-H and not C(sp²)-H bonds.

The contacts from H(C2') and H(C4') (Table 3) are of the order of the sum of the van der Waals radii of H

and O. The distances and angles involved (Table 3) indicated some attractive interaction between the C(sp³)-H and the potential hydrogen-bond acceptors.

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The Structure of *p*-Dimethylaminobenzoic Acid

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Abstract. C₉H₁₁NO₂, triclinic, $P\bar{1}$, $a = 8.598$ (7), $b = 8.061$ (7), $c = 6.652$ (8) Å; $\alpha = 66.099$ (6), $\beta = 94.62$ (5), $\gamma = 90.97$ (5)°, $D_m = 1.29$, $D_x = 1.31$ g cm⁻³, $Z = 2$. The structure was determined from three-dimensional X-ray diffraction data and refined by the least-squares method to $R = 0.113$. The dimethyl-

amino and carboxyl groups are displaced slightly from the plane of the benzene ring and the N atom is non-planar. Pairs of molecules link together to form dimers through two O-H···O hydrogen bonds (2.622 Å) and the N atom is free from any binding of either type N-H···O or N-H···C.