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## Structure of 3'-O-Acetylthymidine

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Abstract.  $C_{12}H_{16}N_2O_6$ ,  $M_r = 284.3$ , orthorhombic,  $P2_{1}2_{1}2_{1}$ , a = 5.485 (4), b = 14.043 (7), c = 17.145 (12) Å, U = 1320.6 Å<sup>3</sup>, Z = 4,  $D_x = 1.43$  g cm<sup>-3</sup>, Mo Ka,  $\lambda = 0.71069$  Å,  $\mu = 1.08$  cm<sup>-1</sup>, F(000) = 600, T = 293 K, R = 0.049 for 1769 observed reflections. The molecule adopts an *anti* conformation, with a sugar pucker of  ${}^{2}T_{3}$  [C(2')*endo*/C(3')-*exo*].

Introduction. The structure determination was performed as part of a series of studies of acetylated nucleosides intended to investigate base-pairing patterns, propeller twisting and acetyl conformations in such compounds. Earlier work has revealed possible biologically interesting acetyl conformations (Wilson & Tollin, 1985) and induced base-pairing in such substituted nucleosides (Wilson, 1985). Since both features of acetyl-base stacking and base-pairing are present in the structure of the closely related compound 3',5'-di-O-acetylthymidine (Wilson, Low, Tollin & Wilson, 1984), it was hoped that the title compound would also show these features.

**Experimental.** The crystals used were grown from aqueous solution and that used in the structural determination had dimensions  $1.2 \times 0.34 \times 0.19$  mm. Cell dimensions and orientation were obtained from the 254, 082, 0,0,10 and symmetry-related reflections. Data were collected on a Nicolet P3 four-circle diffractometer in Aberdeen. 2168 reflections were measured using Mo Ka radiation. Data collection

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parameters:  $0 < 2\theta < 60^{\circ}$ ,  $\theta/2\theta$  scans with a  $2\theta$  scan range of  $2.41-2.8^{\circ}$  depending on  $2\theta$  value, and a variable scan rate in the range  $5.33^{\circ}$  min<sup>-1</sup> (I < 150) to  $58.6^{\circ}$  min<sup>-1</sup> (I > 2500). Range of indices:  $h \le 7$ ,  $k \le 19$ ,  $l \le 24$ . The reflections 284 and 082 were measured every 50 reflections and showed no significant change in their intensities. No absorption or extinction corrections were applied.

Structure solution and refinement. The Pattersonmethods program PATMET (Wilson & Tollin, 1986) was used to obtain an oriented model of the planar base group in the structure which was used as *ab initio* structural knowledge (Main, 1976) in the directmethods package MITHRIL (Gilmore, 1984). A direct-methods run led to the appearance of most non-hydrogen atoms in the first calculated E map. Fourier recycling completed the definition of the structure, and a series of difference Fourier maps revealed all the H-atom positions. All non-hydrogen atoms were refined anisotropically and the H atoms isotropically. In the final stages of refinement all H parameters were fixed. Refinement was by least squares on F using the BLOCK program in the GX system (Mallinson & Muir, 1985). Final refinement parameters: R = 0.049 for 1769 observed reflections  $[I > 3\sigma(I)]; wR = 0.053, w = 1/[\sigma(F)]^2; 181 \text{ param-}$ eters refined; max. shift/e.s.d. = 0.022; max., min. difference peaks = 0.22 and -0.27 e Å<sup>-3</sup>. Scattering factors taken from International Tables for X-ray Crystallography (1974). The PLUTO program (Motherwell & Clegg, 1978) was used for structure drawing. All other calculations were carried out using the GX package. All computations were performed on

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the Rutherford Appleton Laboratory Neutron Division HUB computer, a VAX 8600.

**Discussion.** The atomic numbering used is shown in the perspective drawing (Fig. 1) with atomic parameters, bond lengths and angles given in Tables 1 and 2.\*

The only noteworthy feature in the bond parameters is that the ring exocyclic carbonyl bonds are somewhat longer [C(2)-O(2), 1.205(4); C(4)-O(4),1.234 (4) Å] than the acetyl carbonyl bond [C(3'1)– O(3'1), 1.188 (6) Å]. The short value for this latter bond length is in agreement with other acetyl carbonyl bond lengths in this series of compounds (Wilson, 1985). The length of these short carbonyl bonds seems not to be related to the thermal vibration parameters on either the C or O of the carbonyl group (Wilson & Tollin, 1988).

The pyrimidine base and the acetyl group are approximately planar, as expected, with the largest deviations from the mean planes being atom N(1)[0.03 (1) Å] in the base and atom O(3') [0.02 (1) Å] in the acetyl group.

Molecular conformation. The molecule adopts an anti conformation about the N-glycosidic bond, with  $\chi$ [C(2)-N(1)-C(1')-O(4')] = -123.5 (4)°, and exhibits a sugar pucker of  ${}^{2}T_{3}$  [C(2')-endo/C(3')-exo] described by the pseudorotation parameters P =173.9 (5)°,  $\psi_m = 26.0 \ (5)^\circ.$ The C(4')-C(5')conformation is gg (gauche-gauche), with  $\gamma$ [O(5')- $C(5')-C(4')-C(3') = 50.1 (4)^{\circ}$ . All of these conformational parameters are quite normal for a pyrimidine nucleoside.

Hydrogen bonding. Unlike the closely related 3',5'di-O-acetylthymidine (Wilson, Low, Tollin & Wilson, 1984) the structure does not exhibit base-pairing, there

\* Lists of structure factors, anisotropic thermal parameters, H-atom parameters and a full list of bond lengths and angles have been deposited with the British Library Document Supply Centre as Supplementary Publication No. SUP 44744 (17 pp.). Copies may be obtained through The Executive Secretary, International Union of Crystallography, 5 Abbey Square, Chester CH1 2HU, England.

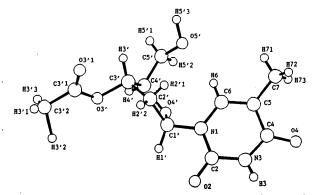


Fig. 1. Perspective view of the title compound, showing atomic numbering.

Table 1. Fractional atomic coordinates and equivalent isotropic temperature factors  $(Å^2)$ , with e.s.d.'s in the least significant digits in parentheses

$U_{\rm eq} = \frac{1}{3}(U_{11} + U_{22} + U_{33}).$					
	x	у	Ζ	$U_{eq}$	
N(1)	-0.7310 (5)	-0.3528 (2)	0.2117 (1)	0.030 (2)	
C(2)	-0.5469 (6)	-0.2947 (2)	0.1846 (2)	0.027 (2)	
O(2)	-0-3998 (5)	-0.2547 (2)	0-2252 (1)	0.040 (1)	
N(3)	-0.5457 (5)	-0.2850 (2)	0.1037(1)	0.026(1)	
C(4)	-0.7040 (6)	-0.3266 (2)	0.0508 (2)	0.026 (2)	
O(4)	-0.6716 (4)	-0.3122 (2)	-0·0195 (1)	0.032 (1)	
C(5)	-0.8968 (5)	-0.3837 (2)	0.0842 (2)	0.029 (2)	
C(7)	<b>−1·0788 (8)</b>	-0·4310 (3)	0.0319 (2)	0.040 (2)	
C(6)	-0.9006 (6)	-0·3944 (2)	0.1624 (2)	0.030 (2)	
C(1′)	-0.7337 (6)	-0.3784 (2)	0-2951 (2)	0.029 (2)	
C(2′)	<i>—</i> 0∙9651 (7)	<i>—</i> 0∙3464 (2)	0.3380 (2)	0.035 (2)	
C(3')	<b>−1·0259 (6)</b>	0·4279 (3)	0.3920 (2)	0.033 (2)	
O(3′)	-0-9018 (5)	-0·4170 (2)	0-4669 (1)	0.041 (2)	
C(3'1)	1·0191 (8)	-0.3632 (2)	0-5212 (2)	0.042 (3)	
O(3'1)	<i>−</i> 1·2104 (6)	<i>—</i> 0∙3259 (2)	0.5099 (2)	0.059 (2)	
C(3'2)	-0·8770 (11)	-0·3619 (3)	0.5956 (2)	0.068 (4)	
C(4′)	-0-9147 (7)	0-5155 (2)	0.3533 (2)	0.033 (2)	
C(5')	-1·0956 (8)	<i>−</i> 0∙5784 (3)	0-3113 (2)	0.045 (2)	
O(5′)	-1.2472 (5)	-0.5265 (2)	0.2599 (2)	0.049 (2)	
O(4′)	-0.7319 (5)	-0-4805 (2)	0-2995 (1)	0.034 (1)	

Table 2. Interatomic distances (Å) and angles (°), excluding H atoms

380 (5)	N(1) C(6) = 1	A A 4 4 1
	N(1) - C(0) = 1	·386 (5)
		·205 (4)
		·386 (5)
		.445 (5)
497 (5)	C(5)-C(6) 1	-349 (5)
534 (5)		.436 (4)
510 (5)	C(3') = O(3') 1	•461 (4)
525 (5)	O(3') - C(3'1) = 1	.361 (5)
188 (6)	C(3'1)-C(3'2) 1	-495 (6)
512 (6)	C(4')-O(4') 1	•449 (5)
414 (5)		
122.4 (3)	C(2) = N(1) = C(1')	118-6 (3)
		124.8 (3)
		121.8 (3)
		118.7 (3)
		125.7 (3)
		117.9 (3)
		123.2 (3)
		107.0 (3)
105-9 (3)	C(1') - C(2') - C(3')	104-8 (3)
110.9 (3)	C(2') - C(3') - C(4')	104.8 (3)
106.3 (3)	C(3') = O(3') = C(3'1)	116-1 (3)
123-3 (4)	O(3')-C(3'1)-C(3'2	) 110.2 (4)
126.5 (4)	C(3') - C(4') - C(5')	114.6 (4)
106-3 (3)	C(5') - C(4') - O(4')	110.4 (3)
112.5 (3)	C(1') - O(4') - C(4')	111.5 (3)
	475 (4) 392 (4) 234 (4) 497 (5) 534 (5) 510 (5) 525 (5) 188 (6) 512 (6) 414 (5) 122-4 (3) 113-3 (3) 127-5 (3) 115-6 (3) 119-8 (3) 122-3 (4) 113-7 (3) 105-9 (3) 110-9 (3) 106-3 (3) 126-5 (4) 106-3 (3)	$\begin{array}{rrrrrrrrrrrrrrrrrrrrrrrrrrrrrrrrrrrr$

being only one hydrogen bond present between the bases. This is between O(4) and N(3) and is detailed as follows:

O(4)...N(3) 
$$(-0.5 + x, -0.5 - y, -z)$$
 2.857 (3) Å  
O(4)...H(3)  $(-0.5 + x, -0.5 - y, -z)$  1.96 (2) Å

The molecules pack in the crystal structure as shown in Fig. 2, with lines along c stacked in a head-to-head and tail-to-tail arrangement, with the head-to-head molecules linked by the hydrogen bond. The molecules

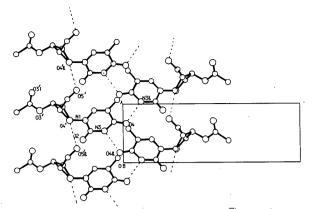


Fig. 2. The stacking in the title compound, showing the hydrogenbonding scheme.

are linked along a by a hydrogen bond between O(4')and O(5'):

O(4')····O(5') 
$$(1 + x, y, z)$$
 2·819 (4) Å  
O(4')····H(5'3)  $(1 + x, y, z)$  2·21 (3) Å.

Thus there are regions of bases and of sugar groups packing along a, as shown.

There are no other significant close contacts between the molecules in the structure. This goes against the strong trend in other acetylated pyrimidine and purine nucleosides for the C=O moiety of an acetyl group to stack above the base ring of a neighbouring molecule (Wilson, 1985). However, the interaction in 3'.5'-di-O-acetvlthymidine was the weakest of those found in the earlier work and while the 3'-acetyl group was involved it is not particularly surprising that in the less sterically congested mono-substituted title compound the interaction no longer plays a significant role. It has been found that in nucleoside structures the mono-acetylated compounds show less propensity to display this interaction.

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# Ammonium Chloride Complexes with 18-Crown-6

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Abstract. (I):  $2C_{12}H_{24}O_{6}\cdot 2NH_{4}^{+}\cdot Cl^{-}\cdot PF_{6}^{-}\cdot 0\cdot 9CH_{2}Cl_{2}$ ,

with  $|F_{0}| > 3\sigma(|F_{0}|)$ . (II):  $C_{12}H_{24}O_{6}NH_{4}^{+}.Cl^{-}.2H_{2}O_{6}$  $\sum_{i=1}^{2} \sum_{j=1}^{2} \sum_{i=1}^{2} \sum_{i=1}^{2} \sum_{i=1}^{2} \sum_{i=1}^{2} \sum_{i$ 1875 (1) Å<sup>3</sup>, Z = 4,  $D_x = 1.26$  Mg m<sup>-3</sup>,  $\lambda$ (Co Ka) = 1.54178 Å,  $\mu = 2.13$  mm<sup>-1</sup>, F(000) = 768, room

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