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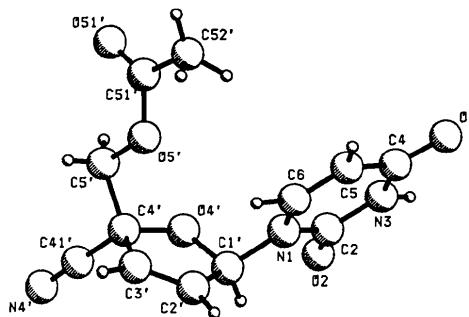


Fig. 1. PLUTO (Motherwell & Clegg, 1978) drawing of the molecule.

*Acta Cryst.* (1992). **C48**, 2277–2278

## Structure of a 4'-C-Branched 2',3'-Didehydro-2',3'-dideoxyuridine

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(Received 29 June 1992; accepted 24 August 1992)

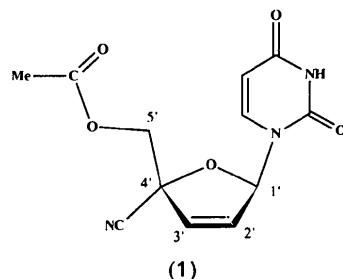
### Abstract

The structure of a 4'-C-branched 2',3'-unsaturated uracil nucleoside, 4'-cyano-2',3'-didehydro-2',3'-dideoxyuridine 5'-acetate, has been determined. The *N*-glycoside torsion angle  $\chi$  has a value of  $-82.7(3)^\circ$  in the *anti* range. The C4'—C5' side-chain conformation is +sc with the torsion angle  $\gamma(C3'—C4'—C5'—O5') = 47.2(4)^\circ$ . The sugar ring is essentially planar. The conformational parameters are in accordance with the IUPAC-IUB Joint Commission on Biochemical Nomenclature [*Pure Appl. Chem.* (1983), **55**, 1273–1280] guidelines.

### Comment

Despite the recent finding that 4'-cyanothymidine acts as a potent inhibitor of the replication of human immunodeficiency virus (O-Yang, Wu, Fraser-Smith & Walker, 1992), only two methods for introducing carbon substituents to the 4'-position of nucleosides have been available. These are 4'-hydroxymethylation of nucleoside 5'-aldehydes (Youssefeyeh, Tegg, Verheyden, Jones & Moffatt, 1977) and a Claisen-type rearrangement of nucleoside 4',5'-enamines (Sechrist & Winter, 1978). A new approach to produce a series of 4'-C-branched 2',3'-unsaturated uracil nucleosides has been disclosed quite recently (Haraguchi, Tanaka, Itoh, Saito & Miyasaka, 1992), wherein carbon

nucleophiles generated from organosilicon reagents of various types can be transferred to a 3',4'-unsaturated derivative through an allylic rearrangement. The use of trimethylsilyl cyanide in this reaction gave an epimeric mixture of two 4'-cyano derivatives from which the title compound (**1**) was isolated as the major product after deprotection followed by acetylation. This molecule shows an *anti* conformation with respect to the sugar ring:  $\chi(C2—N1—C1'—O4') = -82.73(3)^\circ$ . The sugar ring is essentially planar. The maximum displacement from the least-squares plane is 0.023 Å (O4'). The C4'—C5' side-chain conformation is +sc with the torsion angle  $\gamma(C3'—C4'—C5'—O5') = 47.2(4)^\circ$ . Two weak C—H—O intermolecular hydrogen bonds are observed: O2—C1'(1 -  $x$ ,  $1/2 + y$ ,  $2 - z$ ) 3.174(4) Å and O4—C1'( $x$ ,  $y$ ,  $1 + z$ ) 3.148 Å.



### Experimental

#### Crystal data

$C_{12}H_{11}N_3O_5$	Cu $K\alpha_1$ radiation
$M_r = 277.24$	$\lambda = 1.5405 \text{ \AA}$
Monoclinic	Cell parameters from 20 reflections
$P2_1$	$\theta = 29.0\text{--}30.5^\circ$
$a = 14.870 (1) \text{ \AA}$	$\mu = 0.913 \text{ mm}^{-1}$
$b = 5.411 (1) \text{ \AA}$	$T = 297 \text{ K}$
$c = 8.150 (1) \text{ \AA}$	Plate
$\beta = 95.71 (2)^\circ$	$0.55 \times 0.50 \times 0.08 \text{ mm}$
$V = 652.5 (1) \text{ \AA}^3$	Clear
$Z = 2$	
$D_x = 1.411 \text{ Mg m}^{-3}$	

*Data collection*

Rigaku AFC-5 diffractometer  
 $\omega/2\theta$  scans  
 Absorption correction:  
 none  
 1196 measured reflections  
 1165 independent reflections  
 1056 observed reflections  
 $[F > 3\sigma(F)]$

*Refinement*

Refinement on  $F^2$   
 Final  $R = 0.047$   
 $wR = 0.046$   
 $S = 3.04$   
 1056 reflections  
 225 parameters  
 All H-atom parameters refined  
 Calculated weights  
 $w=1/[\sigma^2(F)+0.065F^2]$

$R_{\text{int}} = 0.014$   
 $\theta_{\text{max}} = 60^\circ$   
 $h = -16 \rightarrow 16$   
 $k = 0 \rightarrow 6$   
 $l = 0 \rightarrow 9$   
 3 standard reflections monitored every 150 reflections  
 intensity variation: <3%

( $\Delta/\sigma$ )<sub>max</sub> = 0.164  
 $\Delta\rho_{\text{max}} = 0.228 \text{ e } \text{\AA}^{-3}$   
 $\Delta\rho_{\text{min}} = -0.315 \text{ e } \text{\AA}^{-3}$   
 Atomic scattering factors from *International Tables for X-ray Crystallography* (1974, Vol. IV)

C6—N1—C2	121.7 (2)	C3'—C2'—C1'	110.3 (3)
C6—N1—C1'	122.9 (2)	C2'—C3'—C4'	109.8 (3)
C2—N1—C1'	115.4 (2)	O4'—C4'—C41'	108.0 (3)
O2—C2—N3	122.7 (3)	O4'—C4'—C3'	104.8 (3)
O2—C2—N1	123.2 (2)	O4'—C4'—C5'	109.4 (3)
N3—C2—N1	114.1 (3)	C41'—C4'—C3'	110.8 (3)
C4—N3—C2	127.4 (3)	C41'—C4'—C5'	108.6 (3)
O4—C4—N3	119.4 (4)	C3'—C4'—C5'	115.0 (3)
O4—C4—C5	125.7 (4)	N4'—C41'—C4'	179.5 (5)
N3—C4—C5	114.9 (3)	C4'—O4'—C1'	110.0 (3)
C6—C5—C4	119.6 (4)	O5'—C5'—C4'	106.2 (3)
C5—C6—N1	122.2 (3)	C51'—O5'—C5'	117.2 (3)
O4'—C1'—N1	109.0 (2)	O51'—C51'—O5'	121.9 (4)
O4'—C1'—C2'	104.9 (2)	O51'—C51'—C52'	126.5 (4)
N1—C1'—C2'	114.2 (2)	O5'—C51'—C52'	111.6 (4)

Data collection and cell refinement: *AFD* (Rigaku Corporation, 1985a). Data reduction and structure refinement: *RCRYSTAN* (Rigaku Corporation, 1985b). Program used to solve structure: *SAPI85* (Yao, Zheng, Qian, Han, Gu & Fan, 1985). Molecular graphics: *ACV* (Stardent Computer Inc, 1990). Software used for publication material: *XPACK* (Yamaguchi, 1987). The  $\omega$ -scan width was  $(1.3 + 0.14\tan\theta)^\circ$  and the scan speed was  $32^\circ \text{ min}^{-1}$ . Refinement was by full-matrix least-squares methods.

Lists of structure factors, anisotropic thermal parameters and H-atom coordinates have been deposited with the British Library Document Supply Centre as Supplementary Publication No. SUP 55476 (6 pp.). Copies may be obtained through The Technical Editor, International Union of Crystallography, 5 Abbey Square, Chester CH1 2HU, England. [CIF reference: AS1021]

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Table 1. Fractional atomic coordinates and equivalent isotropic thermal parameters ( $\text{\AA}^2$ )

	$U_{\text{eq}} = \frac{1}{3} \sum_i \sum_j U_{ij} a_i^* a_j^* \mathbf{a}_i \cdot \mathbf{a}_j$
N1	0.6336 (1)
C2	0.5788 (1)
O2	0.5522 (1)
N3	0.5579 (1)
C4	0.5877 (2)
O4	0.5639 (1)
C5	0.6460 (2)
C6	0.6663 (2)
C1'	0.6539 (1)
C2'	0.6847 (2)
C3'	0.7648 (2)
C4'	0.7970 (2)
C41'	0.8046 (2)
N4'	0.8110 (2)
O4'	0.7275 (1)
C5'	0.8864 (2)
O5'	0.8797 (1)
C51'	0.9422 (2)
O51'	1.0015 (2)
C52'	0.9287 (3)
x	0.2698
y	1.1042 (2)
z	0.036 (1)
$U_{\text{eq}}$	

Table 2. Geometric parameters ( $\text{\AA}$ ,  $^\circ$ )

N1—C6	1.370 (4)	C2'—C3'	1.309 (4)
N1—C2	1.392 (4)	C3'—C4'	1.504 (6)
N1—C1'	1.455 (3)	C4'—O4'	1.422 (4)
C2—O2	1.206 (4)	C4'—C41'	1.489 (5)
C2—N3	1.375 (3)	C4'—C5'	1.521 (5)
N3—C4	1.367 (5)	N4'—C41'	1.139 (5)
C4—O4	1.232 (4)	C5'—O5'	1.420 (6)
C4—C5	1.437 (6)	O5'—C51'	1.338 (4)
C5—C6	1.338 (4)	C51'—O51'	1.184 (6)
C1'—O4'	1.431 (4)	C51'—C52'	1.483 (8)
C1'—C2'	1.491 (6)		