

C2—N2—C3—C9	125.1 (2)	C1—C5—C6—C7	98.9 (3)
N2—C3—C4—O2	74.2 (2)	C1—C5—C6—N3	-83.0 (2)
N2—C3—C4—O3	-105.0 (2)		

Table 3. Hydrogen-bond distances (\AA)

Data for normalized H-atom positions are based on bond lengths of O—H = 0.98, N—H = 1.04 and C—H = 1.08 \AA .

$D \cdots A$	$D \cdots A$	$H_{\text{norm}} \cdots A$
OW1···N3	2.775 (3)	1.81
OW1···O2 ⁱ	2.720 (2)	1.75
OW2···OW1	2.744 (3)	1.80
OW2···OW1 ⁱⁱ	2.796 (2)	1.82
N1···O3 ⁱⁱⁱ	2.854 (2)	1.89
N1···OW2 ^{iv}	2.858 (2)	1.84
N1···O3 ^v	3.007 (2)	2.00
N1···O2 ^{vi}	3.111 (2)	2.44
N2···OW ⁱ	2.819 (2)	1.78
N4···OW2 ^v	2.791 (3)	1.76
C1···O1 ⁱ	3.266 (3)	2.46
C5···OW1 ⁱⁱ	3.600 (3)	2.78
C5···N3 ⁱⁱⁱ	3.384 (3)	2.66
C7···O3 ^{iv}	3.481 (3)	2.62
C8···O1 ^{vi}	3.609 (3)	2.80

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

Data collection: *CAD-4 Software* (Enraf–Nonius, 1989). Cell refinement: *CAD-4 Software*. Data reduction: *CAD-4 Software*. Program(s) used to solve structure: *SHELXS86* (Sheldrick, 1985). Program(s) used to refine structure: *SHELXL93* (Sheldrick, 1993). Molecular graphics: *ORTEPII* (Johnson, 1976). Software used to prepare material for publication: *SHELXL93*.

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Lists of structure factors, anisotropic displacement parameters, H-atom coordinates and complete geometry have been deposited with the IUCr (Reference: HA1164). Copies may be obtained through The Managing Editor, International Union of Crystallography, 5 Abbey Square, Chester CH1 2HU, England.

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α -5-Adamantyl-4'-thio-2'-deoxyuridine Methanol Solvate

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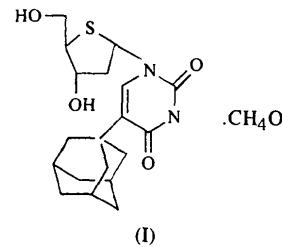
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Abstract

The thiosugar ring of the title compound, $C_{19}H_{26}N_2O_4S \cdot CH_4O$, has the *C*2'-*endo*, *C*3'-*exo* conformation. The orientation of the *C*4'—*C*5' bond is *trans-gauche*, and the glycosidic torsion angle *S*4'—*C*1'—*N*1—*C*6 is $-26.2(2)^\circ$ (*anti*).

Comment

5-Substituted 2'-deoxy-4'-thiouridines are a relatively new group of nucleosides, possessing significant antiviral activity (Dyson, Coe & Walker, 1991; Sechrist, Tiwari, Riordan & Montgomery, 1991). Syntheses provide mixtures of α - and β -anomers, of which only the β -anomers manifest antiviral activity (Rahim *et al.*, 1996). In order to provide information about structure–activity relationships in both anomeric series, the crystal structure of the title compound (**I**), synthesized by Basnak, Sun, Coe & Walker (1996), is presented here. As far as we are aware, it is the first reported crystal structure of an α -anomer within the 2'-deoxy-4'-thiouridine series.



A perspective view of the molecule with atomic numbering is shown in Fig. 1. Bond lengths are normal. The *C*1'—*S*4' and *C*4'—*S*4' bonds are 1.811 (2) and 1.830 (2) \AA , in good agreement with accepted values (Allen *et al.*, 1987). The other bond lengths generally agree well with those found in the crystal structures of 2'-deoxyuridine (Rahman & Wilson, 1972) and α -5-acetyl-2'-deoxyuridine (Hamor, O'Leary & Walker, 1977). The thiosugar ring has the *C*2'-*endo*, *C*3'-*exo* conformation (south), these two atoms being displaced by 0.224 (3) and 0.440 (3) \AA , respectively, on opposite

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sides of the C1'—S4'—C4' plane. The sugar ring of 2'-deoxyuridine has a similar shape; in contrast, the deoxyribose conformation in α -5-acetyl-2'-deoxyuridine is C2'-*endo* (2E). The conformation about C4'—C5' is *trans-gauche*; the C3'—C4'—C5'—O5' torsion angle is $-65.5(2)^\circ$. The glycosidic torsion angle S4'—C1'—N1—C6 is $-26.2(2)^\circ$ (*anti* conformation), similar to the values found in 2'-deoxyuridine (Rahman & Wilson, 1972) and α -5-acetyl-2'-deoxyuridine (Hamor *et al.*, 1977). The adamantyl group has normal bond lengths and angles (C—C bonds 1.516 to 1.544, mean 1.531 Å; C—C—C angles 107.6 to 110.7, mean 109.5°).

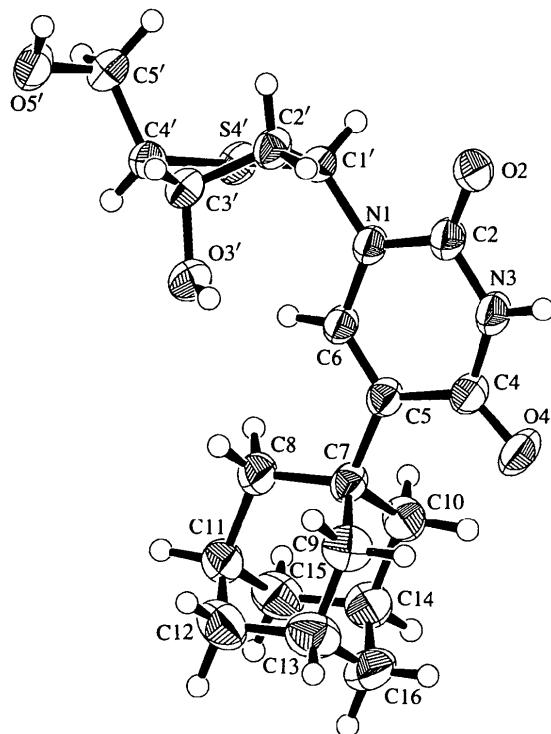


Fig. 1. Molecular structure showing 50% probability displacement ellipsoids (ORTEPII; Johnson, 1976).

In the crystal, nucleoside and methanol molecules are linked through a three-dimensional network of hydrogen bonds. Atoms O3' and O5' of the thiosugar each donate a proton, forming bonds with, respectively, O2 of the pyrimidine base at $-\frac{1}{2} - x, -1 - y, z - \frac{1}{2}$ and O7 of the methanol molecule at $-\frac{3}{2} - x, -1 - y, \frac{1}{2} + z$. Relevant distances and angles are O3'—O2 2.729 (2), O5'—O7 2.852 (2) Å, O3'—H—O2 161, O5'—H—O7 149°. The methanol O atom in turn donates a proton to O4 of a different nucleoside molecule at $x - 1, y, z$ [O7—O4 2.783 (3) Å, O7—H—O4 164°]. Finally, N3 donates a proton to O5' of the thiosugar at $1 + x, y, z$ [N3—O5' 2.839 (2) Å, N3—H—O5' 171°]. Thus, all H atoms attached to the electronegative groups are involved in hydrogen-bond formation, consistent with the principle of maximum hydrogen bonding (Robertson, 1953).

Experimental

The title compound was recrystallized from MeOH–Et₂O.

Crystal data

C ₁₉ H ₂₆ N ₂ O ₄ S·CH ₄ O	Mo K α radiation
M _r = 410.52	$\lambda = 0.71069$ Å
Orthorhombic	Cell parameters from 12001
P ₂ 12 ₁ 2 ₁	reflections (post-refined
<i>a</i> = 10.5520 (10) Å	using complete data set)
<i>b</i> = 24.9930 (18) Å	$\theta = 2.5\text{--}25.6^\circ$
<i>c</i> = 7.9700 (10) Å	$\mu = 0.187$ mm ⁻¹
<i>V</i> = 2101.9 (3) Å ³	<i>T</i> = 293 (2) K
<i>Z</i> = 4	Irregular
<i>D</i> _x = 1.297 Mg m ⁻³	0.5 × 0.4 × 0.4 mm
<i>D</i> _m not measured	Colourless

Data collection

Rigaku R-Axis II area detector	3588 observed reflections [<i>I</i> > 2σ(<i>I</i>)]
Image-plate scans	<i>R</i> _{int} = 0.027
Absorption correction: none	$\theta_{\max} = 25.59^\circ$
12001 measured reflections	$h = -11 \rightarrow 12$
3648 independent reflections	$k = -29 \rightarrow 30$
[<i>I</i> > σ(<i>I</i>)]	$l = -9 \rightarrow 9$

Refinement

Refinement on <i>F</i> ²	$\Delta\rho_{\max} = 0.167$ e Å ⁻³
<i>R</i> [<i>F</i> ² > 2σ(<i>F</i> ²)] = 0.0345	$\Delta\rho_{\min} = -0.227$ e Å ⁻³
<i>wR</i> (<i>F</i> ²) = 0.1049	Extinction correction: none
<i>S</i> = 1.049	Atomic scattering factors
3648 reflections	from International Tables for Crystallography (1992, Vol. C, Tables 4.2.6.8 and 6.1.1.4)
364 parameters	Absolute configuration:
H atoms: see text	Flack (1983)
<i>w</i> = 1/[σ ² (<i>F</i> _o ²) + (0.0708 <i>P</i>) ² + 0.2780 <i>P</i>] where <i>P</i> = (<i>F</i> _o ² + 2 <i>F</i> _c ²)/3	Flack parameter = 0.00 (7)
(Δ/ <i>σ</i>) _{max} = 0.089	

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

	<i>x</i>	<i>y</i>	<i>z</i>	<i>U</i> _{eq}
S4'	-0.53245 (5)	-0.38812 (2)	0.37649 (6)	0.0432 (2)
O2	-0.1775 (2)	-0.47984 (7)	0.4351 (2)	0.0627 (4)
O3'	-0.50508 (14)	-0.44942 (6)	0.0330 (2)	0.0495 (4)
O4	0.03313 (14)	-0.38513 (6)	0.0393 (2)	0.0601 (4)
O5'	-0.83791 (13)	-0.48140 (6)	0.2836 (2)	0.0531 (4)
O7	-0.7716 (2)	-0.41508 (8)	-0.1759 (3)	0.0818 (6)
N1	-0.2904 (2)	-0.41958 (6)	0.2820 (2)	0.0391 (4)
N3	-0.0764 (2)	-0.43131 (7)	0.2339 (2)	0.0480 (4)
C1'	-0.4057 (2)	-0.43699 (7)	0.3775 (2)	0.0388 (4)
C2'	-0.4631 (2)	-0.48851 (7)	0.3070 (3)	0.0426 (4)
C2	-0.1805 (2)	-0.44559 (8)	0.3238 (3)	0.0453 (4)
C3'	-0.5618 (2)	-0.47275 (7)	0.1777 (2)	0.0406 (4)
C4'	-0.6433 (2)	-0.42975 (7)	0.2577 (2)	0.0381 (4)
C4	-0.0699 (2)	-0.39397 (7)	0.1039 (3)	0.0436 (4)
C5'	-0.7455 (2)	-0.45087 (8)	0.3744 (3)	0.0489 (5)
C5	-0.1906 (2)	-0.36949 (7)	0.0597 (2)	0.0395 (4)
C6	-0.2925 (2)	-0.38355 (7)	0.1511 (2)	0.0387 (4)
C7	-0.1982 (2)	-0.33025 (7)	-0.0865 (2)	0.0388 (4)
C8	-0.3350 (2)	-0.31076 (8)	-0.1170 (3)	0.0512 (5)

C9	-0.1515 (2)	-0.35676 (9)	-0.2497 (3)	0.0559 (5)
C10	-0.1160 (2)	-0.28055 (8)	-0.0493 (3)	0.0523 (5)
C11	-0.3396 (2)	-0.27130 (9)	-0.2650 (3)	0.0556 (5)
C12	-0.2925 (3)	-0.29833 (10)	-0.4234 (3)	0.0661 (7)
C13	-0.1560 (3)	-0.31685 (10)	-0.3962 (3)	0.0669 (7)
C14	-0.1221 (3)	-0.24143 (9)	0.1969 (3)	0.0615 (6)
C15	-0.2575 (3)	-0.22304 (9)	-0.2267 (3)	0.0619 (6)
C16	-0.0720 (3)	-0.26845 (12)	-0.3541 (4)	0.0743 (8)
C17	-0.7078 (4)	-0.36671 (13)	-0.2000 (4)	0.0858 (9)

Table 2. Selected geometric parameters (\AA , $^\circ$)

S4'—C1'	1.811 (2)	N3—C2	1.359 (3)
S4'—C4'	1.830 (2)	N3—C4	1.396 (2)
O2—C2	1.233 (2)	C1'—C2'	1.530 (3)
O3'—C3'	1.424 (2)	C2'—C3'	1.517 (3)
O4—C4	1.223 (2)	C3'—C4'	1.517 (2)
O5'—C5'	1.434 (3)	C4'—C5'	1.519 (3)
O7—C17	1.397 (4)	C4—C5	1.456 (3)
N1—C2	1.370 (3)	C5—C6	1.346 (3)
N1—C6	1.378 (2)	C5—C7	1.525 (2)
N1—C1'	1.500 (2)		
C1'—S4'—C4'	95.21 (8)	O3'—C3'—C4'	106.77 (14)
C2—N1—C6	120.5 (2)	C2'—C3'—C4'	106.7 (2)
C2—N1—C1'	115.19 (14)	C3'—C4'—C5'	114.4 (2)
C6—N1—C1'	124.09 (14)	C3'—C4'—S4'	104.94 (12)
C2—N3—C4	127.4 (2)	C5'—C4'—S4'	109.56 (14)
N1—C1'—C2'	112.3 (2)	O4—C4—N3	118.5 (2)
N1—C1'—S4'	113.65 (12)	O4—C4—C5	126.9 (2)
C2'—C1'—S4'	105.88 (13)	N3—C4—C5	114.7 (2)
C3'—C2'—C1'	107.62 (14)	O5'—C5'—C4'	111.0 (2)
O2—C2—N3	122.8 (2)	C6—C5—C4	117.3 (2)
O2—C2—N1	121.7 (2)	C6—C5—C7	122.6 (2)
N3—C2—N1	115.5 (2)	C4—C5—C7	120.1 (2)
O3'—C3'—C2'	111.6 (2)	C5—C6—N1	124.6 (2)
C2—N1—C1'—C2'			-80.6 (2)
C6—N1—C1'—C2'			94.0 (2)
C2—N1—C1'—S4'			159.24 (14)
C6—N1—C1'—S4'			-26.2 (2)
C4'—S4'—C1'—C2'			-8.74 (14)
S4'—C1'—C2'—C3'			33.1 (2)
C1'—C2'—C3'—C4'			-47.8 (2)
C2'—C3'—C4'—S4'			39.2 (2)
C1'—S4'—C4'—C3'			-17.45 (13)
C3'—C4'—C5'—O5'			-65.5 (2)
S4'—C4'—C5'—O5'			176.94 (12)

Image-plate scans were recorded covering 180° of rotation in 3° frames about one axis, with crystal-detector distance 78 mm and exposure time 10 min frame $^{-1}$. The resulting data contains *ca* 86% of the theoretically accessible independent reflections. Coordinates and anisotropic displacement parameters were refined for non-H atoms. H atoms, except those of the methanol methyl group, were located from a difference map and were refined with isotropic displacement parameters. The methanol methyl H atoms were placed in calculated positions and only their isotropic displacement parameters were refined.

Data collection: *R-Axis II Software* (Rigaku Corporation, 1994). Cell refinement: *R-Axis II Software*. Data reduction: *TEXSAN* (Molecular Structure Corporation, 1993). Program(s) used to solve structure: *TEXSAN*. Program(s) used to refine structure: *SHELXL93* (Sheldrick, 1993). Molecular graphics: *ORTEPII* (Johnson, 1976). Software used to prepare material for publication: *SHELXL93*.

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Lists of structure factors, anisotropic displacement parameters, H-atom coordinates and complete geometry have been deposited with the IUCr (Reference: HA1160). Copies may be obtained through The Managing Editor, International Union of Crystallography, 5 Abbey Square, Chester CH1 2HU, England.

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(2aS,3S,6S,7S,7bR)-7-[Dimethylphenyl]-silyl]-2-oxo-6-[(1R,2S)-2-phenylcyclohexyloxy]-2a,3,6,7,7a,7b-hexahydro-2H-1,4,5-trioxa-4a-azacyclopenta[cd]indene-3-carboxylic Acid 1-Methylethyl Ester

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Abstract

The structure of the title compound, $C_{31}H_{39}NO_7Si$, was determined and found to be a fused tricyclic nitroso acetal. Remarkable features include a twist-boat