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## Key indicators

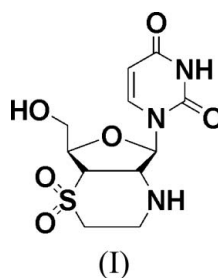
Single-crystal X-ray study  
 $T = 285$  K  
Mean  $\sigma(\text{C}-\text{C}) = 0.004$  Å  
 $R$  factor = 0.041  
 $wR$  factor = 0.108  
Data-to-parameter ratio = 14.9For details of how these key indicators were  
automatically derived from the article, see  
<http://journals.iucr.org/e>.1-[(4*aS*,5*R*,7*R*,7*aS*)-7-Hydroxymethyl-  
3,4,4*a*,5,7,7*a*-hexahydro-2*H*-furo[3,4-*b*]-  
[1,4]thiazin-5-yl]pyrimidine-2,4(1*H*,3*H*)-dione

The title compound,  $\text{C}_{11}\text{H}_{15}\text{N}_3\text{O}_6\text{S}$ , is a 2',3'-thiazine-fused bicyclic nucleoside. The furanose ring adopts a 3'-*endo*,4'-*exo* conformation  ${}_4T^3$ . The orientation of the pyrimidine ring is *anti* with respect to the sugar group. The crystal packing is stabilized by intermolecular N—H...O hydrogen bonds.

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## Comment

A large number of 2',3'-modified nucleosides have been synthesized for the evaluation of their biological activities, revealing that the conformational equilibrium of the sugar unit is a key factor responsible for biological effects (Meldgaard & Wengel, 2000; Len & Mackenzie, 2006). The synthesis of 2',3'-fused bicyclic nucleosides, the analogues conformationally restricted on the sugar ring, has been used as a tool to probe the conformational requirements of key enzymes in order to maximize the therapeutic index of nucleoside anti-viral agents (Vanheusden *et al.*, 2004; Sharma *et al.*, 2004). Here we report the crystal structure of a new bicyclic nucleoside, (I), recently synthesized in our laboratory. Compound, (I), is a 2',3'- $\alpha$ -fused tetrahydrothiazine nucleoside synthesized *via* intramolecular Michael addition of an amino nucleophile to vinyl sulfone.



In (I) (Fig. 1), the five-membered ribose ring C7/C3/C4/C5/O4 adopts an approximate  ${}_4T^3$  conformation, with a pseudorotational phase angle ( $P$ ) of  $36.9(2)^\circ$  and puckering amplitude ( $\tau^m$ ) of  $49.9(3)^\circ$  (Sanger, 1984; Altona & Sundaralingam, 1972; Altona & Sundaralingam, 1973). The glycosidic torsion angle  $\chi$  of  $-160.2(2)^\circ$  shows the orientation of the pyrimidine ring to be *anti* with respect to the sugar group. The torsion angle  $\gamma$  (C4—C5—C6—O3) is  $51.9(4)^\circ$ . The C4—S1 and C3—N1 bond lengths are 1.776(3) and 1.442(4) Å, respectively. The fused six-membered tetrahydrothiazine ring adopts a chair conformation. The crystal packing is stabilized by intermolecular N—H...O hydrogen bonds (Table 1).

The absolute stereochemistry of atoms C3, C4, C5 and C7 has been unambiguously determined to be *S*, *S*, *R* and *R*, respectively, by refining the Flack (1983) parameter.

## Experimental

The title compound was synthesized from 1-[5'-hydroxy-2',3'-di-deoxy-3'-(2-aminoethylsulfonyl)- $\beta$ -D-glyceropent-2'-enofuranosyl]-uracil *via* intramolecular Michael addition (details will be published elsewhere). It was crystallized slowly from a mixture of ethanol and methanol (1:1) at 298 K.

### Crystal data

$C_{11}H_{15}N_3O_6S$   
 $M_r = 317.32$   
 Monoclinic,  $P2_1$   
 $a = 5.365$  (4) Å  
 $b = 7.130$  (5) Å  
 $c = 17.272$  (14) Å  
 $\beta = 90.39$  (3)°  
 $V = 660.7$  (9) Å<sup>3</sup>

$Z = 2$   
 $D_x = 1.595$  Mg m<sup>-3</sup>  
 Mo  $K\alpha$  radiation  
 $\mu = 0.28$  mm<sup>-1</sup>  
 $T = 285$  (2) K  
 Block, yellow  
 $0.34 \times 0.23 \times 0.14$  mm

### Data collection

Rigaku R-Axis RAPID  
 diffractometer  
 $\omega$  scans  
 Absorption correction: multi-scan  
 (ABSCOR; Higashi, 1995)  
 $T_{\min} = 0.917$ ,  $T_{\max} = 0.966$

6286 measured reflections  
 2893 independent reflections  
 2342 reflections with  $I > 2\sigma(I)$   
 $R_{\text{int}} = 0.032$   
 $\theta_{\text{max}} = 27.5^\circ$

### Refinement

Refinement on  $F^2$   
 $R[F^2 > 2\sigma(F^2)] = 0.041$   
 $wR(F^2) = 0.108$   
 $S = 1.08$   
 2893 reflections  
 194 parameters  
 H atoms treated by a mixture of  
 independent and constrained  
 refinement

$w = 1/[\sigma^2(F_o^2) + (0.0595P)^2]$   
 where  $P = (F_o^2 + 2F_c^2)/3$   
 $(\Delta/\sigma)_{\text{max}} < 0.001$   
 $\Delta\rho_{\text{max}} = 0.32$  e Å<sup>-3</sup>  
 $\Delta\rho_{\text{min}} = -0.27$  e Å<sup>-3</sup>  
 Absolute structure: Flack (1983),  
 1263 Friedel pairs  
 Flack parameter: 0.00 (9)

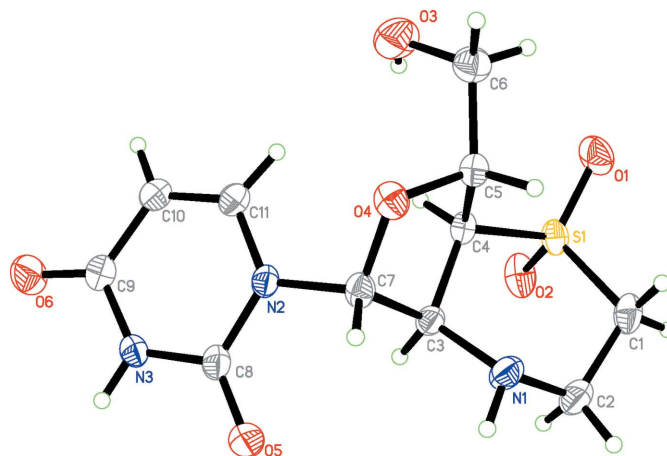
**Table 1**

Hydrogen-bond geometry (Å, °).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
$N3-H3A\cdots O6^i$	0.86	1.92	2.769 (4)	171
$N1-H1\cdots O3^{ii}$	1.00 (4)	2.58 (4)	3.410 (5)	141 (3)
$O3-H3B\cdots O5^{iii}$	0.82	2.69	3.112 (4)	114

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

The C-bound H atoms were positioned geometrically ( $C-H = 0.93-0.98$  Å) and treated as riding on their parent atoms, with  $U_{\text{iso}}(H) = 1.2U_{\text{eq}}(C)$ . The hydroxy H atom was treated as riding on its parent atom, with  $O-H = 0.82$  Å and  $U_{\text{iso}}(H) = 1.5U_{\text{eq}}(O)$ . H3A was positioned geometrically and allowed to ride on N3, with  $U_{\text{iso}}(H) =$



**Figure 1**

The molecular structure of (I) showing the atom-labelling scheme. Displacement ellipsoids are drawn at the 30% probability level. H atoms are represented by circles of arbitrary radius.

$1.2U_{\text{eq}}(N3)$ . H1 was located in a difference Fourier map and refined freely, with  $U_{\text{iso}}(H) = 1.2U_{\text{eq}}(N)$ .

Data collection: *RAPID-AUTO* (Rigaku, 1998); cell refinement: *RAPID-AUTO*; data reduction: *CrystalStructure* (Rigaku, 2002); program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *PLATON* (Spek, 2003); software used to prepare material for publication: *SHELXL97*.

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