

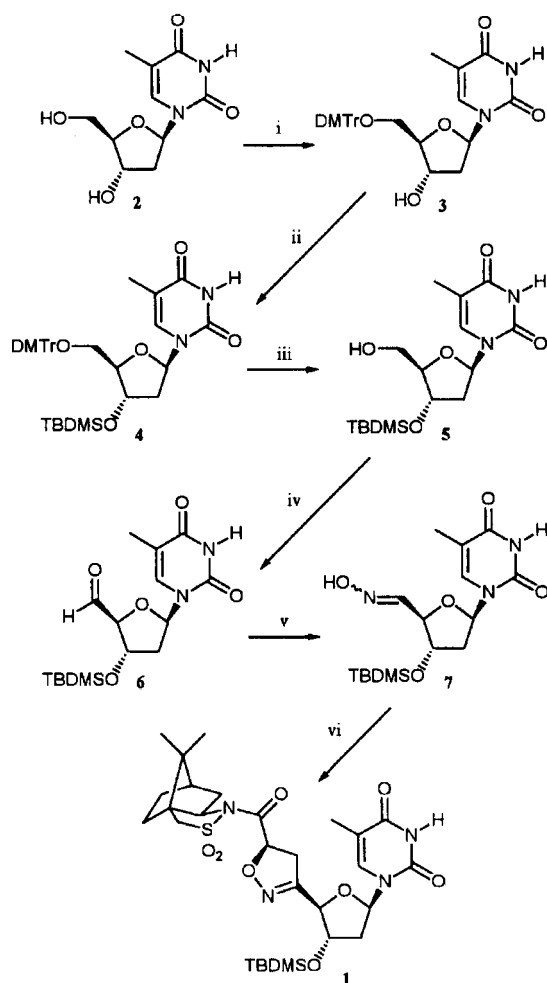
Synthesis and X-Ray Crystal Structure of a Thymidine Derivative

Su Jeong Kim, Ju Young Lee, Ki Hwan Ko, and Byeang Hyeon Kim*

Department of Chemistry, Center for Biofunctional Molecules, Pohang University of Science and Technology, Pohang 790-784, Korea

Received April 25, 1998

There have been numerous research activities in antisense oligonucleotides for the treatment of diseases at the level of gene expression.¹ For the antisense oligonucleotides with heterocyclic isoxazoline linkage,² we synthesized thymidine derivative **1** from thymidine as shown in Scheme 1. Selective protection of 5'-hydroxyl group with 4,4'-dimethoxytrityl (DMTr) group followed by silylation of 3'-hydroxyl group with *t*-butyl dimethyl silyl chloride provided the fully protected compound **4**. Selective deprotection of 5'-position with ZnBr₂, and the modified Moffatt oxidation³



Scheme 1. Reagents and conditions: i, DMTrCl, triethylamine, DMAP, Py, 93%; ii, TBDMSCl, diisopropylethylamine, DMF, 97%; iii, ZnBr₂, CHCl₃/MeOH (9:1), 89%; iv, EDC, DMSO, Py, Trifluoroacetic acid; v, NH₂OH·HCl, Na₂CO₃, MeOH/H₂O (1:1), 70% (iv and v, overall); vi, 1) 0 °C, NaOCl, N-acryloyl (2*R*)-bornane-10,2-sultam, CH₂Cl₂, 2) CH₃SCH₃, 3) recrystallization, 72%.

with 1-(3-dimethyl-aminopropyl)-3-ethylcarbodiimide hydrochloride (EDC) gave the desired aldehyde **6**. This aldehyde was further reacted with hydroxylamine hydrochloride to give the mixture of *syn* and *anti* oxime **7**. Diastereoselective cycloaddition of *in situ* generated nitrile oxide from the oxime **7** to *N*-acryloyl (2*R*)-bornane-10,2-sultam afforded 90:10 mixture of compound **1** and its isoxazoline diastereomer.⁴ It was essential to treat the crude cycloadducts with methyl sulfide to improve the yield. Otherwise the *N*-chlorinated side product at thymine base was isolated in *ca.* 25% yield. Recrystallization from dichloromethane/diethylether/*n*-hexane (1:1:1) solution gave an enantiomerically pure compound **1**.

X-ray diffraction-grade single crystals of compound **1** were grown by slow solvent evaporation of an EtOH/EtOAc solution of the thymidine derivative. The resulting X-

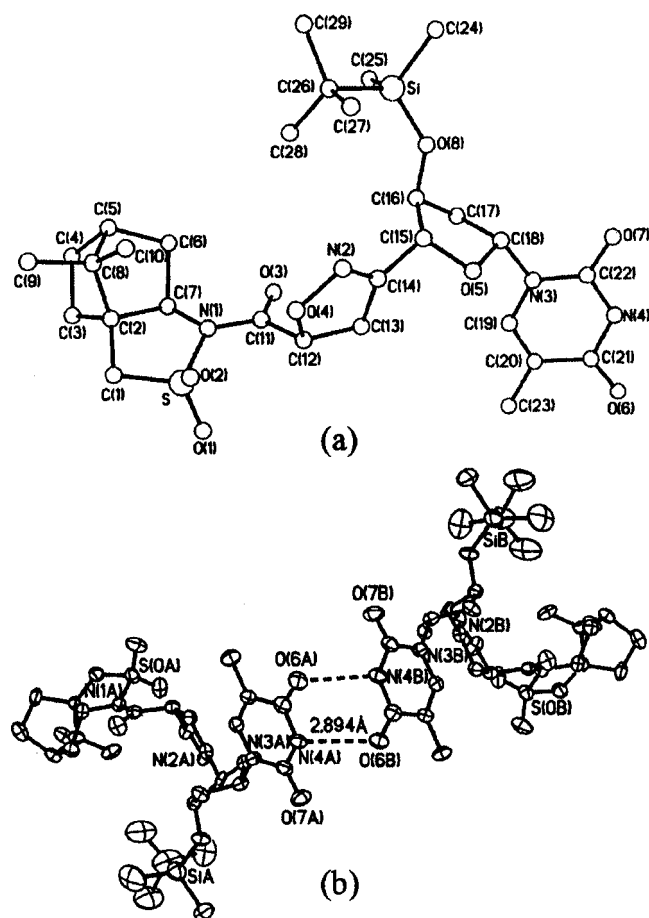


Figure 1. (a) The asymmetric unit of compound **1** with atomic numbering scheme. (b) the dimeric structure of self-assembled **1**.

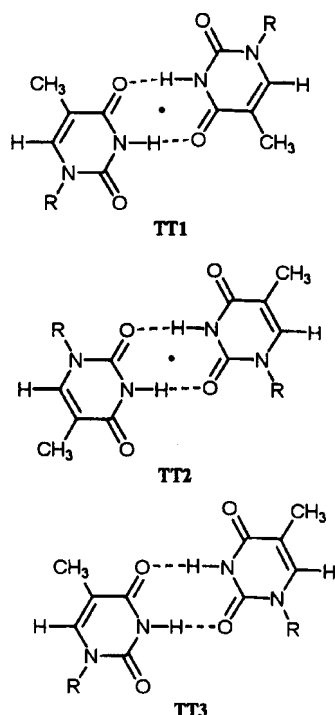


Figure 2. The possible dimeric arrangements of 1-substituted thymines.

ray crystal structure⁵ (Figure 1) reveals a rare example of self-association in the crystal structure of nucleosides.⁶ The number of crystal structures in which self-base pairing occurs is relatively small compared with the number of possibilities. Among the possible dimeric arrangements (Figure 2) of 1-substituted thymine, the **TT1** centrosymmetrical cyclic dimer was mainly formed in the case of 1-methyl thymine.⁷

Molecules of compound **1** are linked *via* N-H...O hydrogen bonds to produce dimers across centers of symmetry (Figure 1(b)). The distance between N and O is 2.894 (0.017) Å, which is slightly longer than the reported values (2.841 Å and 2.830 Å) of 1-methyl thymine⁷ that is closely related to the nucleoside thymidine. As found in most crystal structures involving thymine and uracil derivatives, oxygen O (6) is favored in hydrogen bond formation rather than oxygen O (7). This may be attributable to the higher double bond character of C (22)-O (7) than C (21)-O (6). The preference to centrosymmetrical configuration in thymine-thymine base pairs is due to the antiparallel orientation of the dipole moments.⁶ This results in a favorable cancellation of the total electric field over the crystal volume.

X-ray crystallographic study of thymidine derivative **1** clearly indicates that self-assembly with thymine-thymine base pairing is still applicable in thymidine nucleosides and cyclic dimerization occurs based on the hydrogen bonding between N (4)-H...O (6). However, this type of cyclic self-assembly was not observed in the crystal structure of the thymidine itself.⁸ For thymidine, the carbonyl and N-H groups, which undergo Watson-Crick hydrogen bonding in DNA, are found to hydrogen bonded to 3' or 5' hydroxyl

groups of the sugar moiety. Thus 3' and 5'-hydroxyl groups of sugar play an important role in forming intermolecular hydrogen bonds. Because both 3' and 5'-hydroxyl groups of the sugar are protected in the thymidine derivative **1**, only self-assembly through hydrogen bonding between thymine-thymine base is possible and observed in the crystal structure. For the favorable homo base pairs with centrosymmetrical configuration in thymidine nucleoside, both 3' and 5'-hydroxyl groups should be modified appropriately.

In summary, a thymidine derivative **1** has been synthesized from thymidine in 6 steps (40.5% overall yield) and X-ray crystallographic study of compound **1** shows that this thymidine derivative forms a centrosymmetrical cyclic dimer through hydrogen bonding between thymine-thymine bases.

Acknowledgment. This research was supported by the Ministry of Education (BSR1 97-3437) and POSTECH (1RB9711501). We thank Professor Kimoon Kim and Dr. Dongmok Whang for the X-ray crystallography and helpful comments.

References

- (a) Uhlmann, E.; Peyman, A. *Chem. Rev.* **1990**, *90*, 543. (b) De Mesmaeker, A.; Häner, R.; Martin, P.; Moser, H. E. *Acc. Chem. Res.* **1995**, *28*, 366. (c) Egli, M. *Angew. Chem. Int. Ed. Engl.* **1996**, *35*, 1894.
- Kim, S. J.; Lee, J. Y.; Kim, B. H. *Bioorg. Med. Chem. Lett.* **1998**, *8*, 1313.
- (a) Bousquie, I.; Madiot, V.; Florent, J. C.; Monneret, C. *Bioorg. Med. Chem. Lett.* **1996**, *6*, 1815. (b) Lebreton, J.; De Mesmaeker, A.; Waldner, A.; Fritsch, V.; Wolf, R. M.; Freier, S. M. *Tetrahedron Lett.* **1993**, *34*, 6383.
- (a) Chung, Y. J.; Ryu, E. J.; Keum, G.; Kim, B. H. *Bioorg. Med. Chem.* **1996**, *4*, 209. (b) Kim, B. H.; Curran, D. P. *Tetrahedron* **1993**, *49*, 293. (c) Lee, J. Y.; Kim, B. H. *Tetrahedron* **1996**, *52*, 571. (d) Kim, K. S.; Kim, B. H.; Park, W. M.; Cho, S. J.; Mhin, B. J. *J. Am. Chem. Soc.* **1993**, *115*, 7472.
- Crystal data for 1*: C₂₀H₄₄N₄O₈SSi, M=636.83, crystal system: monoclinic, space group: C2, a=24.491(5) Å, b=7.705(2) Å, c=22.131(4) Å, β=118.52(3)°, V=3520 (1) Å³, Z=4, d_{calc}=1.202 g cm⁻³, T=296 K, Enraf-Nonius CAD4 diffractometer, Mo Kα (λ=0.71073 Å), m=1.75 cm⁻¹. Structure was solved by Patterson method (SHELXS-86). All nonhydrogen atoms were refined anisotropically (SHELXL-93). Final full matrix least squares refinement on F² with all 2341 reflections and 388 variables converged to R1 (I > 2σ(I))=0.096, wR2 (all data)=0.2966 and GOF=1.075.
- Jeffrey, G. A.; Seanger, W. *Hydrogen Bonding in Biological Structures*, Springer-Verlag: Berlin Heidelberg, 1991.
- (a) Kwick, A.; Koetzle, T. F.; Thomas, R. *J. Chem. Phys.* **1974**, *61*, 2711. (b) Hoogsteen, K. *Acta. Cryst.* **1963**, *16*, 28.
- (a) Young, D. W.; Tollin, P.; Wilson, H. R. *Acta. Cryst.* **1969**, *B25*, 1423. (b) Chekhlov, A. N. *J. Struct. Chem.* **1995**, *36*, 155.