

PCR Amplification of 4'-ThioDNA Using 2'-Deoxy-4'-thionucleoside 5'-Triphosphates

Takamitsu Kojima,[†] Kazuhiro Furukawa,[†] Hideto Maruyama,[‡] Naonori Inoue,[‡] Noriko Tarashima,[†] Akira Matsuda,^{*,‡} and Noriaki Minakawa^{*,†}

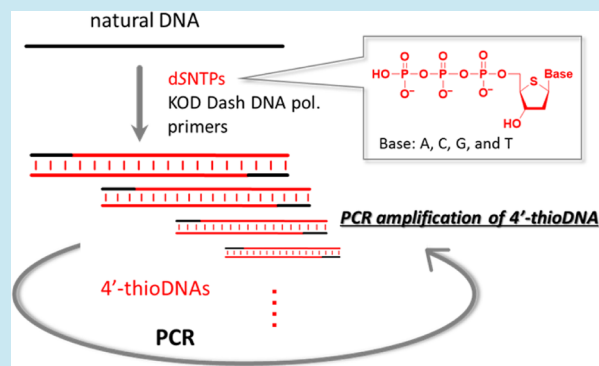
[†]Graduate School of Pharmaceutical Sciences, The University of Tokushima, Shomachi 1-78-1, Tokushima 770-8505, Japan

[‡]Faculty of Pharmaceutical Sciences, Hokkaido University, Kita-12, Nishi-6, Kita-ku, Sapporo 060-0812, Japan

Supporting Information

ABSTRACT: 2'-Deoxy-4'-thioribonucleic acid (4'-thioDNA) having a sulfur atom instead of an oxygen atom in the furanose ring has a nuclease resistance and hybridization ability higher than that of natural DNA. Despite its great potential for various biological applications, a long 4'-thioDNA having all four kinds of 2'-deoxy-4'-thionucleosides has not been reported. In this study, we describe systematic analysis of the incorporation of 2'-deoxy-4'-thionucleoside 5'-triphosphates (dSNTPs) using various DNA polymerases. We found that family B DNA polymerases, which do not have 3'→5' exonuclease activity, could efficiently incorporate dSNTPs via single nucleotide insertion and primer extension. Moreover, 104-mer PCR product was obtained even under the conditions in the presence of all four kinds of dSNTPs when KOD Dash DNA polymerase was used. The resulting PCR product was converted into a natural dsDNA by using PCR with dNTPs, and sequencing of the natural dsDNA revealed that the PCR cycle successfully proceeded without losing the sequence information of the template. To the best of our knowledge, this is the first example of accurate PCR amplification of highly modified DNA in the presence of only unnatural dNTPs.

KEYWORDS: 2'-deoxy-4'-thionucleoside 5'-triphosphate, 4'-thioDNA, PCR amplification



Thus far, a large number of chemically modified nucleoside derivatives have been incorporated into oligonucleotides (ONs) for use in biological, bioengineering, and therapeutic applications. In general, these ONs are prepared via chemical methods using the corresponding phosphoramidite units. This method seems to be the most reliable for incorporating a modified unit into the desired position in the ON sequences. However, it is unsuitable for preparing long chain sequences. There is a possibility that an alternative method involving enzymatic synthesis using the corresponding modified nucleoside 5'-triphosphates (NTPs) could be applied. Since this method affords much longer sequences from readily available natural DNA templates, if successful, it would be a versatile method for use in a variety of applications. Therefore, intense efforts are currently underway toward efficient enzymatic synthesis of modified ONs by using not only primer extension and PCR amplification using DNA polymerases but also *in vitro* transcription with RNA polymerases.^{1–10} A typical example of this research area is the investigation of the evolution of aptamers.^{11–17} To the best of our knowledge, there are very few examples in which PCR proceeds in the presence of all four kinds of chemically modified dNTPs instead of natural dNTPs. Andreola et al. reported PCR in the presence of all four kinds of α -phosphorothioate dNTPs (α -PS-dNTPs); however, the incorporation efficiency of the full length product was very

low.¹⁸ Furthermore, no PCR examination has been successful using all four kinds of sugar-modified dNTP analogues. Quite recently, two groups have individually reported attractive results concerning selection of aptamers. Thus, Yu et al. succeeded to isolate fully modified aptamers consist of (3',2')- α -L-threose nucleic acid (TNA) using all four kinds of TNA triphosphates,^{19,20} while Pinherio et al. isolated fully modified 1,5-anhydrohexitol nucleic acid (HNA) aptamers using all four kinds of HNA triphosphates,²¹ respectively. However, these outstanding results demonstrated were based on a combination of DNA-polymerase-mediated primer extension and reverse-transcription, but not PCR, which is generally used for *in vitro* selection. Therefore, there is no example of a highly modified nucleic acid analogue that can be amplified by direct PCR amplification. Since the resulting fully modified DNA analogues are expected to be synthetic genetic polymers,^{19–21} and also to isolate functionalized aptamers, the development of chemically modified dNTPs, which can be amplified by PCR is highly desirable.

Our group has been developing a series of 4'-thionucleic acids, made up of 4'-thioribonucleosides²² and 2'-deoxy-4'-

Received: June 27, 2013

Published: August 7, 2013

thionucleosides.²³ We expected that the resulting 4'-thionucleic acids, that is, 4'-thioRNA and 4'-thioDNA, would exhibit not only high hybridization abilities and nuclease resistance arising from chemical modification but also biological equivalency with natural RNA and DNA because sulfur and oxygen atoms belong to the same group in the periodic table. In fact, these chemically modified ONs showed such favorable properties.^{24–27} and therefore were utilized in gene silencing via RNAi machinery.^{28–30} In addition, together with these successful results obtained by using a chemical approach, we have reported the isolation of partially modified 4'-thioRNA aptamers by using test tube evolution, since the 4'-thiopyrimidine nucleoside 5'-triphosphates (SUTP and SCTP) were good substrates for *in vitro* transcription by using T7 RNA polymerase.³¹ Concerning 4'-thioDNA, we could amplify a partially modified 4'-thioDNA containing 2'-deoxy-4'-thiothymidine and 2'-deoxy-4'-thiocytidine units by using PCR in the presence of 2'-deoxy-4'-thiothymidine and 2'-deoxy-4'-thiocytidine 5'-triphosphates (dSTTP and dSCTP, Figure 1) with natural purine dNTPs

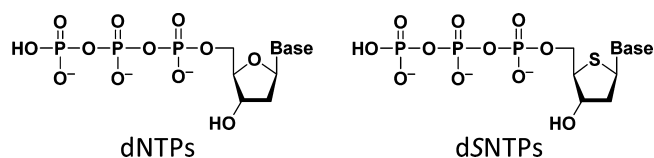


Figure 1. Chemical structures of 2'-deoxyribonucleoside 5'-triphosphates (dNTPs) and 2'-deoxy-4'-thioribonucleoside 5'-triphosphate (dSNTPs). Base = A, C, G, or T.

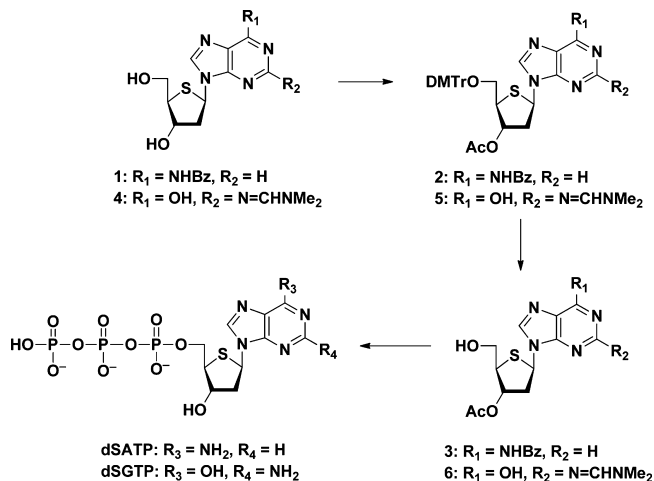
(dATP and dGTP). The amplified partially modified 4'-thioDNAs were used as templates for transcription to give RNA sequences *in vitro* and in mammalian cells.³² This was the first example that 2'-deoxy-4'-thiopyrimidine nucleoside 5'-triphosphates as natural congeners could act as substrates for DNA polymerases and that the resulting partially modified 4'-thioDNAs were highly useful materials (synthesis and enzymatic recognition of 2'-deoxy-5-ethyl-4'-thiouridine 5'-triphosphate, which is not a 4'-S congener of natural dNTPs, are reported as one exception; see ref 33). These observations strongly suggested that 4'-thioDNAs as well as 4'-thioRNAs would be good candidates for not only chemically modified aptamers but also synthetic genetic polymers, if fully modified 4'-thionucleic acids could be freely obtained. Thus, we synthesized additional triphosphates, i.e., dSATP and dSGTP (Figure 1), and investigate their enzymatic recognition along with dSTTP and dSCTP by using DNA polymerases.

In this work, we first screened DNA polymerases by using primer extension in the presence of all four kinds of dSNTPs. As a result, polymerases belonging to family A, such as the Klenow fragment and Taq DNA polymerase, and reverse transcriptases (RTs) did not incorporate dSNTPs efficiently, whereas those belonging to family B, such as Therminator and KOD Dash DNA polymerases, effectively incorporated dSNTPs. Next, using family B DNA polymerases, PCR amplification in the presence of dSNTP(s) was examined, and we finally succeeded in the enzymatic synthesis of a 104-mer 4'-thioDNA by PCR in the presence of all four kinds of dSNTPs. Details of primer extension reactions and PCR amplifications in the presence of dSNTP(s) are described in this paper.

RESULTS AND DISCUSSION

Chemistry. The syntheses of dSATP and dSGTP are shown in Scheme 1. Starting with *N*-benzoyl-2'-deoxy-4'-thioadeno-

Scheme 1. Syntheses of 2'-Deoxy-4'-thioadenosine 5'-Triphosphate (dSATP) and 2'-Deoxy-4'-thioguanosine 5'-Triphosphate (dSGTP)



sine (1),²³ two hydroxyl groups were successively protected with dimethoxytrityl (DMTr) and acetyl groups to give 2. The DMTr group of 2 was then removed by treatment with 80% aqueous AcOH to give 3, a substrate for the subsequent phosphorylation reaction. Following the efficient method reported by Ludwig and Eckstein,³⁴ 3 was treated with 2-chloro-4*H*-1,3,2-benzodioxaphosphorin-4-one, followed by bis-(tri-*n*-butylammonium)pyrophosphate. Then the resulting cyclotriphosphate intermediate was oxidized with 1% iodine in pyridine-H₂O, and successive treatment with NH₄OH gave the desired dSATP in 50% yield after DEAE Sephadex column chromatographic purification. In the same manner, 4 was converted into 6 via fully protected 5, and dSGTP was prepared from 6 by using the one-pot reaction described above.

Primer Extension with dSNTPs Using a Series of DNA Polymerases. We have previously reported that Therminator and KOD Dash DNA polymerases are the best for incorporating dSTTP and dSCTP under PCR conditions.³² To further investigate the differences in the recognition of dSNTPs, we first carried out primer extension assays in the presence of dSNTPs using a series of DNA polymerases (Figure 2). The natural 5'-FITC labeled primer (20-mer) and template I (27-mer) were used in this experiment (Figure 2A). Thirteen DNA polymerases in total were chosen from families A, B, and Y DNA polymerases and reverse transcriptase (RT) for these assays. As shown in Figure 2B, almost no extension was observed when family A DNA polymerases and RT (lanes 2–5, 12 and 13) were used, although Tth DNA polymerase (lane 4) and M-MLV RT (lane 12) gave partial extension products. Dpo4 DNA polymerase (family Y) (lanes 14 and 15) gave better results than family A and RT ones did, although a strong band was observed at the position where three nucleotides were incorporated. On the other hand, the DNA polymerases belonging to family B (lanes 6–11) showed stronger activity than those belonging to families A and Y DNA polymerases or RT did. Of these polymerases, Vent (exo⁻) (lane 6), Deep Vent (exo⁻) (lane 7), Therminator (lane 10), and KOD Dash (lane 11) DNA polymerases showed notable

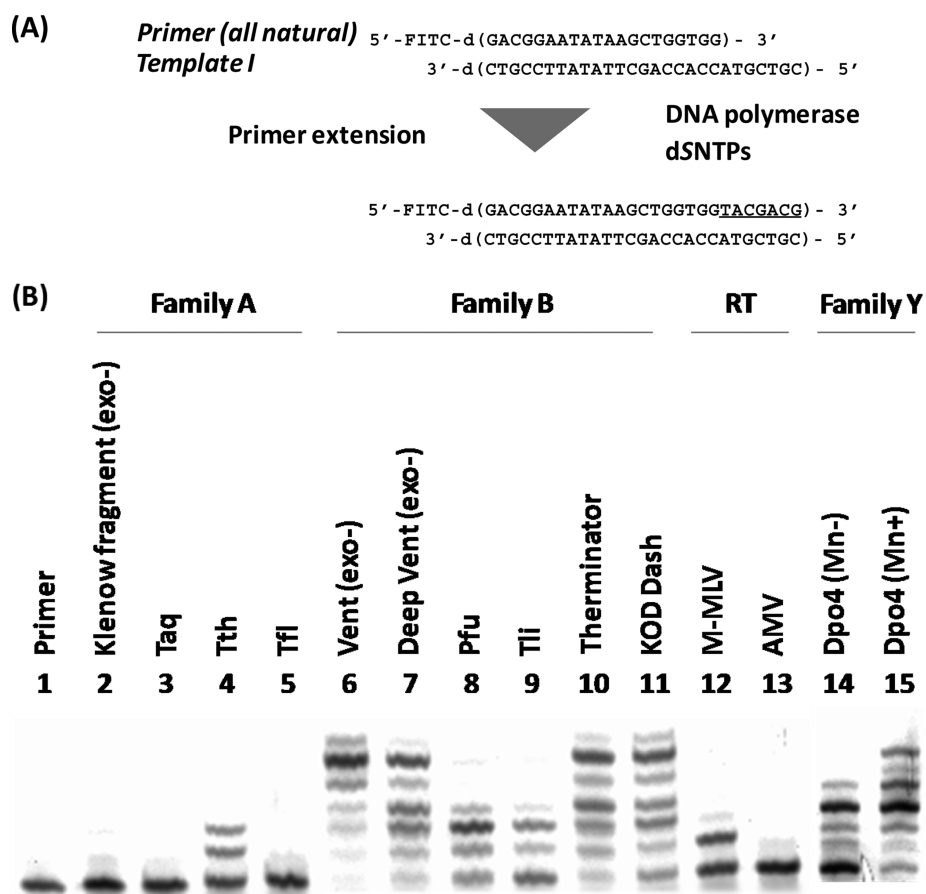


Figure 2. Primer extension by various DNA polymerases with dSNTPs and natural DNA template. (A) Sequences of DNA templates and primer. Underlined are extended nucleotides. (B) Denaturing polyacrylamide gel image of the products. Annealed 5'-FITC labeled primer (0.8 μ M) and template I (1.0 μ M) mixture, dSNTP (40 μ M), and DNA polymerases (0.01–0.4 U/ μ L) in a reaction buffer (25 μ L) were incubated at 37 or 74 $^{\circ}$ C.

efficiency for primer extension. It has been reported that family B DNA polymerases can be utilized to incorporate not only base-modified nucleotides but also sugar-modified nucleotides.^{1,7–17} Our results indicated that family B DNA polymerases were considerably tolerant to dSNTPs. The Pfu (lane 8) and Tli (lane 9) DNA polymerases showed less activity than other family B DNA polymerases did. This is probably due to their higher fidelity to dNTPs caused by 3'→5' exonuclease activity.³⁵ The family Y DNA polymerase Dpo4 (lane 14) showed activity comparable to that of the family B DNA polymerases especially when manganese ion was added (lane 15).³⁶ Family Y DNA polymerases have an ability to incorporate damaged dNTPs, such as 8-hydroxy-dGTP and 2-hydroxy-dATP.³⁷ This low fidelity to dNTPs of Dpo4 DNA polymerase could make it possible to incorporate sugar-modified dNTP analogues such as dSNTPs. From these results, there was a significant difference in the potential to incorporate dSNTPs among the families of DNA polymerases.

Kinetic Studies of the Incorporation of dSNTPs via Single Nucleotide Insertion. From the primer extension assays, we found that the DNA polymerases belonging to family B gave better results than those of families A and Y DNA polymerases or RT did. Next, we carried out single nucleotide insertion assays using Therminator DNA polymerase (family B) or the Klenow fragment (exo⁻) (family A) to further investigate the efficiencies and selectivities of dSNTP incorporation by using steady-state kinetics. Templates I–IV were prepared to investigate each d(S)NTP incorporation

(Table 1). The kinetic parameters (K_m = the Michaelis constant, V_{max} = the maximum rate of the enzyme reaction, and V_{max}/K_m = the insertion efficiency) were determined for each dSNTP with each primer-template set at various concentrations of triphosphates, and then the incorporation efficiencies relative to those of the corresponding natural dNTPs were calculated.

As shown in Table 1, the kinetic parameters for dTTP and dSTTP in the presence of a natural primer were compared. The relative efficiencies of the incorporations of dSNTPs (35–140) by using Therminator were comparable with those of dNTPs (100) (Table 1A). In contrast, the same experiment with the Klenow fragment showed that the relative efficiencies for dSNTPs incorporation were from 1 to 5 since the K_m values for dSNTPs (1.5–12 μ M) were significantly higher than those for dNTPs (0.077–0.15 μ M), as shown in Table 1B. This result indicates that Therminator incorporates dSNTP more efficiently than the Klenow fragment (exo⁻) does because of its higher affinity for dSNTPs.

Li et al. have solved the atomic-level structures of complexes of the Klenow fragment with dNTPs and have revealed that the phenylalanine at position 667 (Phe667) is highly conserved and forms a platform, which is nearly parallel to the sugar rings of the incoming dNTPs.³⁸ In addition, it has been shown that the side chain of Phe667 is critical in distinguishing dNTPs from other sugar motifs as substrates.³⁹ Since the conformation of the sugar ring significantly changes upon substituting 4'-oxygen with 4'-sulfur due to the differences in the atomic radii (oxygen

Table 1. Steady-State Kinetic Parameters for Insertion of Single Nucleotides into Template-Natural Primer Duplexes by Therminator DNA Polymerase (A) and Klenow Fragment (exo⁻) (B)

<i>Primer (natural)</i> 5' -FITC-d (GACGGAATATAAGCTGGTGG) - 3'				
<i>Template I (for d(S)TTP)</i> 3' -d (CTGCCTTATATTGACCACC A TGCTGC) - 5'				
<i>Template II (for d(S)CTP)</i> 3' -d (CTGCCTTATATTGACCACC G TGCTGC) - 5'				
<i>Template III (for d(S)ATP)</i> 3' -d (CTGCCTTATATTGACCACC T TGCTGC) - 5'				
<i>Template IV (for d(S)GTP)</i> 3' -d (CTGCCTTATATTGACCACC C TGCTGC) - 5'				
d(S) NTP	K _m (μM)	V _{max} (% min ⁻¹)	efficiency V _{max} /K _m (% min ⁻¹ M ⁻¹)	relative efficiency
(A) Therminator				
dTTP	0.044 ± 0.03	0.36 ± 0.03	8.1 × 10 ⁶	100
dSTTP	0.011 ± 0.013	0.12 ± 0.03	11 × 10 ⁶	140
dCTP	0.18 ± 0.03	0.28 ± 0.06	1.6 × 10 ⁶	100
dSCTP	0.25 ± 0.01	0.27 ± 0.04	1.1 × 10 ⁶	69
dATP	0.042 ± 0.021	0.24 ± 0.06	5.7 × 10 ⁶	100
dSATP	0.030 ± 0.022	0.21 ± 0.07	7.0 × 10 ⁶	120
dGTP	0.075 ± 0.014	0.38 ± 0.07	5.1 × 10 ⁶	100
dSGTP	0.18 ± 0.04	0.26 ± 0.03	1.8 × 10 ⁶	35
(B) Klenow Fragment (exo ⁻)				
dTTP	0.15 ± 0.06	0.15 ± 0.02	1.0 × 10 ⁶	100
dSTTP	8.5 ± 1.8	0.18 ± 0.03	2.1 × 10 ⁴	2
dCTP	0.077 ± 0.027	0.33 ± 0.05	4.4 × 10 ⁶	100
dSCTP	12 ± 6	0.34 ± 0.07	2.9 × 10 ⁴	1
dATP	0.13 ± 0.05	0.33 ± 0.03	2.5 × 10 ⁶	100
dSATP	1.5 ± 0.9	0.19 ± 0.04	1.3 × 10 ⁵	5
dGTP	0.15 ± 0.05	0.47 ± 0.01	3.1 × 10 ⁶	100
dSGTP	12 ± 5	0.47 ± 0.02	4.1 × 10 ⁴	1

60 pm vs sulfur 100 pm), bond lengths (C–O 1.428 Å vs C–S 1.839 Å), and torsion angles (C–O–C 114.2° vs C–S–C 94.6°), the steric barrier caused by Phe667 could lead to a loss of recognition of dSNTPs by the Klenow fragment (exo⁻). Meanwhile, Yang et al. have studied the structure of RB69 DNA polymerase, which belongs to family B.⁴⁰ They have

shown that the equivalent position of Phe667 in the Klenow fragment (exo⁻) is substituted with asparagine (Asn564). The amide side chain of Asn564 forms a hydrogen bond via a water molecule with the nonbridging oxygen of the β-phosphate of the incoming dNTP, and steric hindrance around the sugar ring of dSNTPs has little effect. Since the asparagine is highly conserved around this site in family B DNA polymerases, it should be possible to incorporate dSNTPs as a substrate without a decrease in the affinities. These results are consistent with reports from other groups, which show that family B DNA polymerases can incorporate sugar-modified nucleoside 5'-triphosphates better than family A ones do.^{1,7,9–17}

As described above, the efficiency of single nucleotide insertion of dSNTPs was comparable to that of natural dNTPs when natural primer was used. In order to further evaluate the efficiency of dSNTP incorporation, we next carried out similar assays using the natural-thio chimera primer and the thio primer (Table 2). The natural-thio chimera primer contained five consecutive 2'-deoxy-4'-thionucleotides at the 3'-end, and the thio primer consisted of only 2'-deoxy-4'-thionucleotides. As a result, the relative efficiencies of dSNTP incorporation significantly decreased when the natural-thio chimera primer was used. Interestingly, the relative efficiencies of dSNTP incorporation improved when the thio primer was used, except in the case of dSCTP. Thus, the relative efficiencies followed the trend of natural primer > thio primer > natural-thio chimera primer. These results suggest that the elongation of dSNTPs by using Therminator is relatively slow in the initial phase and recovers in a later phase after incorporation of several nucleotides (all of the information on steady-state kinetic parameters are presented in Supplementary Table S1).

As described earlier, a large number of sugar-modified nucleoside triphosphates have been prepared, and their single nucleotide insertion, followed by primer extension, has been investigated extensively. In many cases, the relative efficiencies of single nucleotide insertion are comparable to that of natural dNTP when natural primer is used. However, the efficiencies

Table 2. Steady-State Kinetic Parameters for Insertion of Single Nucleotides into Template-Natural, Natural-Thio, and Thio Primer Duplexes by Therminator DNA Polymerase^a

<i>Primer (natural)</i> 5' -FITC-d (GACGGAATATAAGCTGGTGG) - 3'						
<i>Primer (natural-thio)</i> 5' -FITC-d (GACGGAATATAAGCTGGTGG) - 3'						
<i>Primer (thio)</i> 5' -FITC-d (GACGGAATATAAGCTGGTGG) - 3'						
<i>Template I (for d(S)TTP)</i> 3' -d (CTGCCTTATATTGACCACC A TGCTGC) - 5'						
<i>Template II (for d(S)CTP)</i> 3' -d (CTGCCTTATATTGACCACC G TGCTGC) - 5'						
<i>Template III (for d(S)ATP)</i> 3' -d (CTGCCTTATATTGACCACC T TGCTGC) - 5'						
<i>Template IV (for d(S)GTP)</i> 3' -d (CTGCCTTATATTGACCACC C TGCTGC) - 5'						
dSNTP	primer	K _m (μM)	V _{max} (% min ⁻¹)	efficiency V _{max} /K _m (% min ⁻¹ M ⁻¹)	Relative efficiency	
dSTTP	natural	0.011 ± 0.013	0.12 ± 0.03	11 × 10 ⁶	100	
	natural-thio	0.11 ± 0.09	0.025 ± 0.010	2.2 × 10 ⁵	2	
	thio	0.092 ± 0.036	0.25 ± 0.01	2.7 × 10 ⁶	25	
dSCTP	natural	0.25 ± 0.01	0.27 ± 0.04	1.1 × 10 ⁶	100	
	natural-thio	0.034 ± 0.017	0.014 ± 0.002	4.0 × 10 ⁵	36	
	thio	1.7 ± 0.12	0.35 ± 0.01	2.1 × 10 ⁵	20	
dSATP	natural	0.030 ± 0.022	0.21 ± 0.07	7.0 × 10 ⁶	100	
	natural-thio	0.12 ± 0.07	0.021 ± 0.005	1.8 × 10 ⁵	3	
	thio	0.041 ± 0.016	0.15 ± 0.01	3.6 × 10 ⁶	51	
dSGTP	natural	0.18 ± 0.04	0.26 ± 0.03	1.8 × 10 ⁶	100	
	natural-thio	0.054 ± 0.017	0.043 ± 0.019	8.0 × 10 ⁵	44	
	thio	0.067 ± 0.001	0.11 ± 0.00	1.7 × 10 ⁶	94	

^aUnderlined sequences indicate 4'-thio nucleotides.

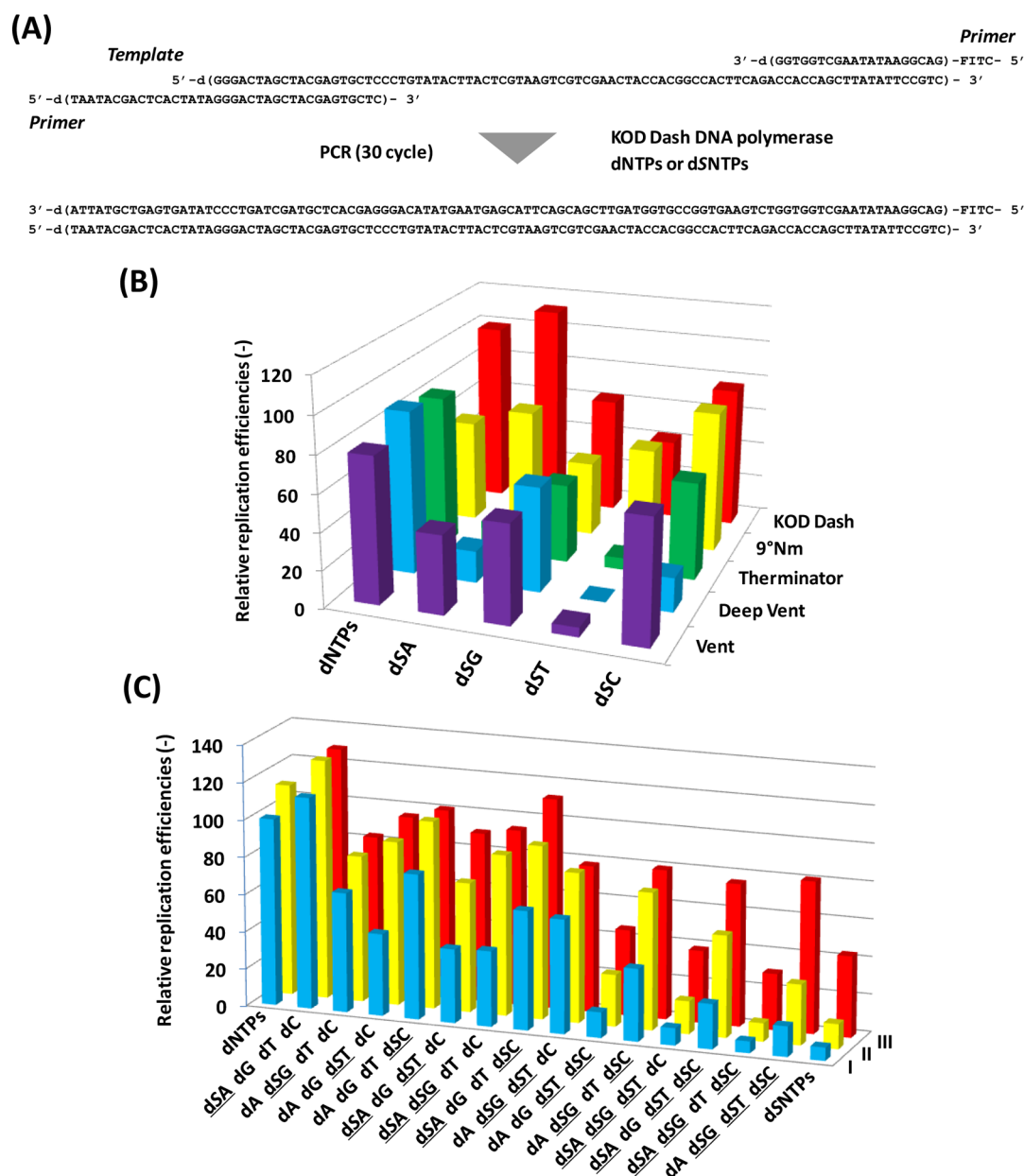


Figure 3. PCR in the presence of dSNTPs. (A) Sequences of template and primers for PCR. (B) Comparison of the yields of PCR products using KOD Dash, Vent, Deep Vent, Terminator, 9°Nm, and Pfu DNA polymerase in the presence of all natural dNTPs or dNTPs containing one kind of dSNTPs. Y-axis indicates the replication yields relative to that of KOD Dash DNA polymerase in the presence of natural dNTPs. (C) Relative replication efficiencies of PCR. Y-axis indicates the replication yields relative to that with natural dNTPs in case of condition I. Reactions were performed in 20 μ L of KOD Dash buffer, 0.5 μ M primers, 0.2 mM corresponding d(S)NTPs, and 0.1 pmol of DNA template containing 1 U of KOD Dash DNA polymerase (I), 2 U of KOD Dash DNA polymerase (II), or 2 U of KOD dash DNA polymerase and 2% DMSO (III). The extension time for the elongation in PCR was 30 s (I and II) or 10 min (III).

are remarkably lower if the primer already includes modified nucleotide unit(s) on its 3'-end. This is similar to our single nucleotide insertion using a natural-thio chimera primer, and these data agree with those of other sugar-modified nucleoside triphosphates.^{1,10} It is not clear why the relative efficiencies of dSNTP incorporation are higher when a thio primer is used. Further investigations to explain these results are needed and are being performed.

Optimization of the Conditions for the PCR with dSNTPs. We next studied PCR amplification of 4'-thioDNA containing all four kinds of 2'-deoxy-4'-thionucleosides in the presence of dSNTP(s). In our previous communication,³² we have reported the PCR amplification with dSTTP and dSCTP

in the presence of dATP and dGTP using KOD Dash DNA polymerase. The PCR amplification to give partially modified 4'-thioDNAs under the conventional conditions (condition I: primer extension at 72 °C for 30 s, 1 U of KOD Dash DNA polymerase) in the presence of one kind of dSNTP using a variety of PCR enzymes was further investigated with additional dSATP and dSGTP. The template DNA and primers used in this experiment were natural oligonucleotides. The sequences of the 87-mer template and primers used to afford the 104-mer amplified dsDNA are shown in Figure 3A. The PCR products were quantified by measuring the fluorescence intensities of the corresponding bands. When Vent, Deep Vent, or Terminator DNA polymerase was used, PCR was inefficient in the presence

of dSTTP. On the other hand, 9°Nm or KOD Dash DNA polymerases afforded the desired 4'-thioDNA regardless of the conditions (Figure 3B). Since KOD Dash DNA polymerase afforded the best results in the above experiments, further conditions, i.e., in the presence of more than two dSNTPs, were examined. The amount of PCR products remarkably decreased with an increase in the ratio of dSNTPs (Figure 3C, condition I). In order to improve PCR efficiency, we first increased the amount of the polymerase. As a result, some improvement was observed when the amount of the enzyme was increased from 1 U to 2 U (condition II: primer extension at 72 °C for 30 s, 2 U of KOD Dash DNA polymerase). However, the incorporations were insufficient especially in the case where all dNTPs were substituted to dSNTPs even under condition II. Therefore, we tested the addition of DMSO⁴¹ or betaine,⁴² which are known to be effective additives for PCR, or Mn²⁺, which is known to increase the tolerance of polymerases for the incorporation of nucleotides with modifications in the furanose ring, to the reaction mixtures.⁴³ However, no significant improvement was observed when betaine or Mn²⁺ was added to the reaction mixture, whereas the addition of DMSO slightly improved the incorporation of dSNTPs. Accordingly, we increased the extension time for PCR from 30 s to 10 min to improve the incorporation of dSNTPs. Finally, the optimum conditions for PCR in the presence of dSNTPs were determined to be the following: primer extension at 72 °C for 10 min, 2 U of KOD Dash DNA polymerase, and 2% DMSO in the reaction mixture (condition III). The yield of the full-length PCR product was 43.3% when all four kinds of dSNTPs were used under condition III relative to that when only dNTPs were used under condition I (Figure 3C and Supplementary Figure S1; 13.4% under condition I and 7.1% under condition II, respectively).

In order to check the accuracy of the PCR in the presence of dSNTPs, the resulting PCR products from all four kinds of dSNTPs were sequenced. The modified PCR products were first converted into natural dsDNA by using PCR with dNTPs because 4'-thioDNA could not be directly subjected to the sequencing reaction (Supplementary Figure S2). We found that the dSNTPs were incorporated during the PCR cycle without losing the sequence information of the template. As described earlier, there is only one example in which PCR has been carried out in the presence of all four modified dNTPs (α -PS-dNTPs).¹⁸ In that experiment, however, the sequence analysis of the resulting highly modified PCR product showed that there was a large deletion in the central region of the amplified fragment, indicating that incorporation of phosphorothioate dNTPs with a phosphorothioate template was inefficient. Accordingly, the results of PCR experiments is of great value because this is the first report on PCR amplification of unnatural oligomer using all four kinds of unnatural triphosphates with high accuracy.

Conclusion. In summary, we systematically analyzed the incorporation of dSNTPs using DNA polymerases. Primer extension assays using a natural DNA template indicated that family B DNA polymerases effectively incorporated dSNTPs, while families A and Y DNA polymerases and RT did not. Using KOD dash DNA polymerase, sufficient amounts of PCR product, that is, a highly modified 4'-thioDNA, was obtained even in the presence of only dSNTPs by optimizing the conditions. Sequence analysis of the resulting PCR product after conversion into the natural dsDNA revealed that replication of 4'-thioDNA took place with a high accuracy.

Since this is the first example of accurate PCR amplification of highly modified DNA in the presence of only unnatural dNTPs, the resulting 4'-thioDNA would be a good candidate for not only chemically modified aptamers but also synthetic genetic polymers.

METHODS

Synthesis of dSNTPs. All information on the organic synthesis is in the Supporting Information; analytical data of dSATP and dSGTP follows:

2'-Deoxy-4'-thioadenosine 5'-Triphosphate (dSATP). ¹H NMR (D₂O) δ 8.68 (s, 1 H), 8.28 (s, 1 H), 6.30 (t, 1 H, $J = 6.6$ Hz), 4.83 (m, 1 H), 4.35–4.23 (m, 2 H), 3.80 (m, 1 H), 2.74 (m, 2 H); ³¹P NMR (D₂O) δ -6.3, -11.0, -21.8; ESI-LRMS m/z 505.9608 [M-H]⁻; ESI-HRMS calcd for C₁₀H₁₅N₅O₁₁P₃S [M-H]⁻; 505.9702, found 505.9697.

2'-Deoxy-4'-thioguanosine 5'-Triphosphate (dSGTP). ¹H NMR (D₂O) δ 8.21 (s, 1 H), 6.13 (t, 1 H, $J = 6.5$ Hz), 4.80 (m, 1 H), 4.29 (m, 2 H), 3.76 (m, 1 H), 2.75 (m, 1 H), 2.63 (m, 1 H); ³¹P NMR (D₂O) δ -5.3, -10.6, -19.9; ESI-LRMS m/z 521.9624 [M-H]⁻; ESI-HRMS calcd for C₁₀H₁₅N₅O₁₂P₃S [M-H]⁻; 521.9651, found 521.9624.

Primer Extension. For the screening of DNA polymerase, the following enzymes were used: Taq (Takara), M-MLV (Takara), KOD Dash (TOYOBO), Vent (exo⁻) (NEB), Deep Vent (exo⁻) (NEB), Therminator (NEB), Klenow fragment (exo⁻) (Promega), Tth (Promega), Tfl (Promega), Tli (Promega), Pfu (Promega), AMV (Promega), and Dpo4 (Trevigen). Annealed 5'-FITC labeled primer (0.8 μ M) and template I (1.0 μ M) mixture, dSNTP (40 μ M), and DNA polymerases (0.01–0.4 U/ μ L) in a reaction buffer (25 μ L) were incubated at 37 or 74 °C. After 10 min, 10- μ L aliquots of the reaction mixture were sampled and added into the loading solution (10 μ L) containing 10 M urea, 5 mM EDTA, 0.05% (w/v) bromophenol blue, and 0.05% (w/v) xylene cyanol. The samples were analyzed by 20% denaturing polyacrylamide gel electrophoresis (PAGE) containing 8 M urea, which was visualized and analyzed with a FLA-2000 (FUJIFILM).

Single Nucleotide Insertion and Steady-State Kinetics. Insertion reactions were initiated by adding dNTP (0.01–10 μ M) or dSNTP (0.1–5 μ M) to the reaction mixture containing a mixture of duplex consisting of 5'-FITC-labeled primer (0.8 μ M) and DNA template I–IV (1.0 μ M), 0.02 U/ μ L of the Klenow Fragment (exo⁻) in a buffer containing 50 mM Tris-HCl (pH 7.2), 10 mM MgSO₄, and 0.1 mM DDT, or 0.003 U/ μ L of Therminator DNA polymerase in a ThermoPol buffer. Reaction time was adjusted for each combination to give extents of reaction of 25% or less. The reaction was quenched by adding 8 μ L of the stop buffer. The extents were analyzed by 20% denaturing PAGE containing 8 M urea, which was quantitated by a FLA-2000 (FUJIFILM). Reaction velocities were calculated as the yield of reaction divided by reaction time. The kinetic parameters (K_m and V_{max}) were determined by linear regression analysis of a Hanes–Woolf plot with an average of three independent experiments.

PCR with dSNTP. For the first screening of DNA polymerase for PCR, KOD Dash, Vent (exo⁻), Deep Vent (exo⁻), Therminator, and 9°Nm were used. The PCR was performed in 20 μ L of corresponding buffer containing DNA polymerase (0.05 U/ μ L), 87-mer DNA template (5.0 nM), 200 μ M each of dNTPs or dSNTPs, and 0.5 μ M primers. KOD DNA polymerase was utilized for the further optimization. The PCR mixture was prepared in a total volume of 20 μ L by

adding 2 μL of 10 \times KOD DNA polymerase buffer, KOD Dash DNA polymerase (condition I ; 0.05 U/ μL , conditions II and III ; 0.10 U/ μL), 87-mer DNA template (5.0 nM), 200 μM each of dNTPs or dSNTPs, 0.5 μM primers, and 2% DMSO (only in condition III). The reaction mixtures were gently vortexed and then amplified using a thermal cycler (TaKaRa). PCR was performed under the condition I, II [94 $^{\circ}\text{C}$, 15 s; 94 $^{\circ}\text{C}$, 15 s; 50 $^{\circ}\text{C}$, 15 s; 72 $^{\circ}\text{C}$, 30 s] \times 30 cycles; 72 $^{\circ}\text{C}$, 5 min] or condition III [94 $^{\circ}\text{C}$, 15 s; 94 $^{\circ}\text{C}$, 15 s; 50 $^{\circ}\text{C}$, 15 s; 72 $^{\circ}\text{C}$, 10 min] \times 30 cycles; 72 $^{\circ}\text{C}$, 5 min]. PCR products were analyzed by 6.4% PAGE and were quantified by measuring the fluorescence intensities of the corresponding bands with a Molecular Imager FX Pro (Bio-Rad). Prior to a second PCR with dNTPs, the products were purified with High Pure PCR Product Purification Kit (Roche). After the second PCR with dNTPs, the products were repurified with the kit (Roche) and then were sequenced with each of PCR primer by a ABI PRISM 3100 Genetic Analyzer.

■ ASSOCIATED CONTENT

Supporting Information

This material is available free of charge via the Internet at <http://pubs.acs.org>.

■ AUTHOR INFORMATION

Corresponding Author

*Phone: +81-11-706-3228. Fax: +81-11-706-4980. E-mail: matuda@pharm.hokudai.ac.jp (A.M.). Phone: +81-88-633-7299. Fax: +81-88-633-7299. E-mail: minakawa@tokushima-u.ac.jp (N.M.).

Notes

The authors declare no competing financial interest.

■ ACKNOWLEDGMENTS

This work was supported by a Grant-in-Aid for Scientific Research (B) (Grant no. 24390027). We would like to thank Mr. H. Kitaike (Center for Instrumental Analysis, The University of Tokushima) and Ms. M. Kikuchi and Ms. S. Oka (Center for Instrumental Analysis, Hokkaido University) for technical supports of analytical data.

■ REFERENCES

- (1) Heuberger, B. D., and Switzer, C. (2008) A pre-RNA candidate revisited: both enantiomers of flexible nucleoside triphosphates are DNA polymerase substrates. *J. Am. Chem. Soc.* 130, 412–413.
- (2) Eoff, R. L., McGrath, C. E., Maddukuri, L., Salamanca-Pinzon, S. G., Marquez, V. E., Marnett, L. J., Guengerich, F. P., and Egli, M. (2010) Selective modulation of DNA polymerase activity by fixed-conformation nucleoside analogues. *Angew. Chem., Int. Ed.* 49, 7481–7485.
- (3) Ogino, T., Sato, K., and Matsuda, A. (2010) Incorporation of 2'-deoxy-2'-isonucleoside 5'-triphosphates (iNTPs) into DNA by A- and B-family DNA polymerases with different recognition mechanisms. *ChemBioChem* 11, 2597–2605.
- (4) Siegmund, V., Santner, T., Micura, R., and Marx, A. (2011) Enzymatic synthesis of 2'-methylseleno-modified RNA. *Chem. Sci.* 2, 2224–2231.
- (5) Rao, H., Sawant, A. A., Tanpure, A. A., and Srivatsan, S. G. (2012) Posttranscriptional chemical functionalization of azide-modified oligoribonucleotides by bioorthogonal click and Staudinger reactions. *Chem. Commun.* 48, 498–500.
- (6) Carrasco, N., and Huang, Z. (2004) Enzymatic synthesis of phosphoroselenoate DNA using thymidine 5'-(α -P-seleno)-

triphosphate and DNA polymerase for X-ray crystallography via MAD. *J. Am. Chem. Soc.* 126, 448–449.

- (7) Kuwahara, M., Obika, S., Nagashima, J., Ohta, Y., Suto, Y., Ozaki, H., Sawai, H., and Imanishi, T. (2008) Systematic analysis of enzymatic DNA polymerization using oligo-DNA templates and triphosphate analogs involving 2',4'-bridged nucleosides. *Nucleic Acids Res.* 36, 4257–4265.
- (8) Yamashige, R., Kimoto, M., Takezawa, Y., Sato, A., Mitsui, T., Yokoyama, S., and Hirao, I. (2012) Highly specific unnatural base pair systems as a third base pair for PCR amplification. *Nucleic Acids Res.* 40, 2793–2806.
- (9) Tsai, C. H., Chen, J., and Szostak, J. W. (2007) Enzymatic synthesis of DNA on glycerol nucleic acid templates without stable duplex formation between product and template. *Proc. Natl. Acad. Sci. U.S.A.* 104, 14598–14603.
- (10) Chaput, J. C., and Szostak, J. W. (2003) TNA synthesis by DNA polymerases. *J. Am. Chem. Soc.* 125, 9274–9275.
- (11) Ichida, J. K., Zou, K., Horhota, A., Yu, B., McLaughlin, L. W., and Szostak, J. W. (2005) An in vitro selection system for TNA. *J. Am. Chem. Soc.* 127, 2802–2803.
- (12) Peng, C. G., and Damha, M. J. (2007) Polymerase-directed synthesis of 2'-deoxy-2'-fluoro-beta-D-arabinonucleic acids. *J. Am. Chem. Soc.* 129, 5310–5311.
- (13) Veedu, R. N., Vester, B., and Wengel, J. (2008) Polymerase chain reaction and transcription using locked nucleic acid nucleotide triphosphates. *J. Am. Chem. Soc.* 130, 8124–8125.
- (14) Kataoka, M., Kouda, Y., Sato, K., Minakawa, N., and Matsuda, A. (2011) Highly efficient enzymatic synthesis of 3'-deoxyapionucleic acid (apioNA) having the four natural nucleobases. *Chem. Commun.* 47, 8700–8702.
- (15) Dubois, C., Campbell, M. A., Edwards, S. L., Wengel, J., and Veedu, R. N. (2012) Stepping towards highly flexible aptamers: enzymatic recognition studies of unlocked nucleic acid nucleotides. *Chem. Commun.* 48, 5503–5505.
- (16) Nawale, G. N., Gore, K. R., Hobartner, C., and Pradeepkumar, P. I. (2012) Incorporation of 4'-C-aminomethyl-2'-O-methylthymidine into DNA by thermophilic DNA polymerases. *Chem. Commun.* 48, 9619–9621.
- (17) Wheeler, M., Chardon, A., Goubet, A., Morihiro, K., Tsan, S. Y., Edwards, S. L., Kodama, T., Obika, S., and Veedu, R. N. (2012) Synthesis of selenomethylene-locked nucleic acid (SeLNA)-modified oligonucleotides by polymerases. *Chem. Commun.* 48, 11020–11022.
- (18) Andreola, M. L., Calmels, C., Michel, J., Toulme, J. J., and Litvak, S. (2000) Towards the selection of phosphorothioate aptamers optimizing in vitro selection steps with phosphorothioate nucleotides. *Eur. J. Biochem.* 267, 5032–5040.
- (19) Yu, H., Zhang, S., and Chaput, J. C. (2012) Darwinian evolution of an alternative genetic system provides support for TNA as an RNA progenitor. *Nat. Chem.* 4, 183–187.
- (20) Yu, H., Zhang, S., Dunn, M. R., and Chaput, J. C. (2013) An efficient and faithful in vitro replication system for Threose nucleic acid. *J. Am. Chem. Soc.* 135, 3583–3591.
- (21) Pinheiro, V. B., Taylor, A. I., Cozens, C., Abramov, M., Renders, M., Zhang, S., Chaput, J. C., Wengel, J., Peak-Chew, S. Y., McLaughlin, S. H., Herdewijn, P., and Holliger, P. (2012) Synthetic genetic polymers capable of heredity and evolution. *Science* 336, 341–344.
- (22) Naka, T., Minakawa, N., Abe, H., Kaga, D., and Matsuda, A. (2000) The stereoselective synthesis of 4'- β -thioribonucleosides via the Pummerer reaction. *J. Am. Chem. Soc.* 122, 7233–7243.
- (23) Inoue, N., Kaga, D., Minakawa, N., and Matsuda, A. (2005) Practical synthesis of 2'-deoxy-4'-thioribonucleosides: substrates for the synthesis of 4'-thioDNA. *J. Org. Chem.* 70, 8597–8600.
- (24) Hoshika, S., Minakawa, N., and Matsuda, A. (2004) Synthesis and physical and physiological properties of 4'-thioRNA: application to post-modification of RNA aptamer toward NF- κ B. *Nucleic Acids Res.* 32, 3815–3825.
- (25) Inoue, N., Minakawa, N., and Matsuda, A. (2006) Synthesis and properties of 4'-ThioDNA: unexpected RNA-like behavior of 4'-ThioDNA. *Nucleic Acids Res.* 34, 3476–3483.

(26) Matsugami, A., Ohyama, T., Inada, M., Inoue, N., Minakawa, N., Matsuda, A., and Katahira, M. (2008) Unexpected A-form formation of 4'-thioDNA in solution, revealed by NMR, and the implications as to the mechanism of nuclease resistance. *Nucleic Acids Res.* 36, 1805–1812.

(27) Takahashi, M., Minakawa, N., and Matsuda, A. (2009) Synthesis and characterization of 2'-modified-4'-thioRNA: a comprehensive comparison of nuclease stability. *Nucleic Acids Res.* 37, 1353–1362.

(28) Hoshika, S., Minakawa, N., Kamiya, H., Harashima, H., and Matsuda, A. (2005) RNA interference induced by siRNAs modified with 4'-thioribonucleosides in cultured mammalian cells. *FEBS Lett.* 579, 3115–3118.

(29) Hoshika, S., Minakawa, N., Shionoya, A., Imada, K., Ogawa, N., and Matsuda, A. (2007) Study of modification pattern-RNAi activity relationships by using siRNAs modified with 4'-thioribonucleosides. *ChemBioChem* 8, 2133–2138.

(30) Takahashi, M., Nagai, C., Hatakeyama, H., Minakawa, N., Harashima, H., and Matsuda, A. (2012) Intracellular stability of 2'-OMe-4'-thioribonucleoside modified siRNA leads to long-term RNAi effect. *Nucleic Acids Res.* 40, 5787–5793.

(31) Kato, Y., Minakawa, N., Komatsu, Y., Kamiya, H., Ogawa, N., Harashima, H., and Matsuda, A. (2005) New NTP analogs: the synthesis of 4'-thioUTP and 4'-thioCTP and their utility for SELEX. *Nucleic Acids Res.* 33, 2942–2951.

(32) Inoue, N., Shionoya, A., Minakawa, N., Kawakami, A., Ogawa, N., and Matsuda, A. (2007) Amplification of 4'-thioDNA in the presence of 4'-thio-dTTP and 4'-thio-dCTP, and 4'-thioDNA-directed transcription in vitro and in mammalian cells. *J. Am. Chem. Soc.* 129, 15424–15425.

(33) Alexandrova, L. A., Semizarov, D. G., Krayevsky, A. A., and Walker, R. T. (1996) 4'-Thio-5-ethyl-2'-deoxyuridine 5'-triphosphate (TEDUTP): Synthesis and substrate properties in DNA-synthesizing systems. *Antivir. Chem. Chemother.* 7, 237–242.

(34) Ludwig, J., and Eckstein, F. (1989) Rapid and efficient synthesis of 5'-O-(1-thiotriphosphates), 5'-triphosphate and 2',3'-cyclophosphorothioates using 2-chloro-4H-1,3,2-benzodioxaphosphorin-4-one. *J. Org. Chem.* 54, 631–635.

(35) Cline, J., Braman, J. C., and Hogrefe, H. H. (1996) PCR fidelity of pfu DNA polymerase and other thermostable DNA polymerases. *Nucleic Acids Res.* 24, 3546–3551.

(36) Ling, H., Boudsocq, F., Woodgate, R., and Yang, W. (2001) Crystal structure of a Y-family DNA polymerase in action: a mechanism for error-prone and lesion-bypass replication. *Cell* 107, 91–102.

(37) Villani, G., Tanguy Le Gac, N., Wasungu, L., Burnouf, D., Fuchs, R. P., and Boehmer, P. E. (2002) Effect of manganese on in vitro replication of damaged DNA catalyzed by the herpes simplex virus type-1 DNA polymerase. *Nucleic Acids Res.* 30, 3323–3332.

(38) Li, Y., Kong, Y., Korolev, S., and Waksman, G. (1998) Crystal structures of the Klenow fragment of *Thermus aquaticus* DNA polymerase I complexed with deoxyribonucleoside triphosphates. *Protein Sci.* 7, 1116–1123.

(39) Tabor, S., and Richardson, C. C. (1995) A single residue in DNA polymerases of the *Escherichia coli* DNA polymerase I family is critical for distinguishing between deoxy- and dideoxyribonucleotides. *Proc. Natl. Acad. Sci. U.S.A.* 92, 6339–6343.

(40) Yang, G., Franklin, M., Li, J., Lin, T. C., and Konigsberg, W. (2002) Correlation of the kinetics of finger domain mutants in RB69 DNA polymerase with its structure. *Biochemistry* 41, 2526–2534.

(41) Choi, J. S., Kim, J. S., Joe, C. O., Kim, S., Ha, K. S., and Park, Y. M. (1999) Improved cycle sequencing of GC-rich DNA template. *Exp. Mol. Med.* 31, 20–24.

(42) Mytelka, D. S., and Chamberlin, M. J. (1996) Analysis and suppression of DNA polymerase pauses associated with a trinucleotide consensus. *Nucleic Acids Res.* 24, 2774–2781.

(43) Tabor, S., and Richardson, C. C. (1989) Effect of manganese ions on the incorporation of dideoxynucleotides by bacteriophage T7 DNA polymerase and *Escherichia coli* DNA polymerase I. *Proc. Natl. Acad. Sci. U.S.A.* 86, 4076–4080.