Supplementary Information for

Synthesis of activated 3'-amino-3'-deoxy-2-thio-thymidine, a superior substrate for the nonenzymatic copying of nucleic acid templates

Enver Cagri Izgu,^{†,‡} Seung Soo Oh,^{†,‡} and Jack W. Szostak^{†,‡,*}

[†]Howard Hughes Medical Institute, Department of Molecular Biology and Center for Computational and Integrative Biology, Massachusetts General Hospital, 185 Cambridge Street, Boston, Massachusetts 02114, United States

[‡]Department of Genetics, Harvard Medical School, 77 Avenue Louis Pasteur, Boston, Massachusetts 02115, United States

^{*}Department of Chemistry and Chemical Biology, Harvard University, 12 Oxford St., Cambridge, Massachusetts 02138, United States

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1. General Materials and Methods

Pyridine (pyr), dimethylformamide (DMF), trimethyl phosphate $(PO(OMe)_3)$, Phosphorus(V) oxychloride (POCl₃), acetic anhydride and piperidine were purchased as anhydrous grade from Sigma-Aldrich. Ammonia solution (7 N) in methanol, tetramethylguanidine (TMG), 1-(2-hydroxyethyl)-imidazole (HEI), 4-morpholineethane-sulfonic acid sodium salt (MES), 3-(cyclohexylamino)-1-propanesulfonic acid (CAPS) and 4-(2-hydroxyethyl)piperazine-1-ethanesulfonic acid (HEPES) were also purchased from Sigma-Aldrich. A hydrogen sulfide (H₂S) gas tank equipped with an H₂S pressure gauge was purchased from Airgas. Primer (DNA) and template (RNA) oligonucleotide strands were obtained from IDT (Coralville, IA). The primer strand was then enzymatically modified with a 3'-amino-2',3'-dideoxyguanosine triphosphate, which was purschased from TriLink Biotechnologies (San Diego, CA). Silica gel chromatographic purification of the synthesized molecules was performed using Teledyne Isco combiflash instrument equipped with a C18Aq high performance RediSep® column. All primer extension reactions were carried out with solutions prepared in nuclease-free water and at 4 °C. Nuclear Magnetic Resonance (NMR) spectroscopic analyses were carried out on either a Varian INOVA 400 MHz or a Bruker Ascend 400 MHz spectrometer. Chemical shifts (δ) for ¹H NMR spectra were referenced to TMS at $\delta = 0.00$ ppm, to CHD₂OD at $\delta = 3.31$ ppm, to CHD₂S(O)CD₃ at $\delta =$ 2.50 ppm, or to HDO at δ = 4.80 ppm. ¹³C NMR spectra were referenced to CDCl₃ at δ = 77.23 ppm, to CD₃OD at $\delta = 49.15$ ppm, or to CD₃S(O)CD₃ at $\delta = 39.51$ ppm. The ³¹P NMR spectra were referenced externally to P(OMe)₃ (dissolved in CDCl₃) at $\delta = 140.0$ ppm using a co-axial NMR tube. The following abbreviations are used to describe NMR resonances: s (singlet), d (doublet), t (triplet), g (quartet), m (multiplet), br (broad), and nfom (non-first order multiplet). Coupling constants (J) are reported in Hz. Infrared spectra were recorded on an FT-IR spectrometer; the most intense and/or diagnostic peaks are reported. Low-resolution mass spectroscopy (LRMS) analyses were performed using a Bruker Daltonics Esquire 6000 mass spectrometer. Liquid chromatography followed by high-resolution mass spectroscopy (LC-HRMS) analysis in the ESI mode was carried out on an Agilent 6520 Q-TOF LC/MS-MS instrument.

2. Experimental Procedures

1-((2*R*,4*S*,5*S*)-4-azido-5-(hydroxymethyl)tetrahydrofuran-2-yl)-2-ethoxy-5 methyl-pyrimidin-4(1*H*)-one: Compound 3.



To a solution of 2^1 (10 g, 23.8 mmol, 1 equiv) in absolute EtOH (650 mL), was added freshly distilled DBU (20 g, 133.0 mmol, 4 equiv) at room temperature. The reaction mixture was purged with Ar for 15 min and a vertical condenser was attached to the reaction flask, which was then placed in a 90 °C-oil bath. The reaction mixture was stirred for 24 h, then cooled to room temperature. The precipitate was filtered off and the filtrate was concentrated by rotary evaporation. The resulting material was re-dissolved in CH₂Cl₂ (500 mL) washed with water (250 mL) and saturated aqueous NaCl solution (250 mL). The extracted organic layer was dried over Na₂SO₄ and concentrated under reduced pressure. The resulting crude product was purified by combiflash silica-gel chromatography with a CH₂Cl₂-MeOH elution mixture (0-to-10% MeOH gradient) to afford the nucleoside **3** (7.0 g, 81% yield).

¹**H NMR** (400 MHz, CDCl₃) δ 7.99 (s, 1H), 6.15 (t, J = 6.5 Hz, 1H), 4.77 (br m, 1H), 4.48 (q, J = 7.0 Hz, 2H), 4.46 (br dd, J = 5.5, 5.0 Hz, 1H), 4.07–4.04 (br ddd, J = 10.0, 6.0, 3.5 Hz, 1H), 4.03 (dd, J = 4.0, 3.0 Hz, 1H), 3.94–3.89 (br ddd, J = 10.0, 6.0, 3.5 Hz, 1H), 2.46–2.35 (m, 2H), 1.91 (s, 3H), and 1.38 (t, J = 7.0 Hz, 3H); ¹³C NMR (100 MHz, CD₃OD) δ 172.9, 155.0, 134.2, 117.3, 85.9, 85.6, 65.3, 61.7, 60.4, 39.1, 14.4, and 13.9; LRMS (ESI): calcd for C₁₂H₁₇N₅O₄•H⁺ 296.1, found 295.8; calcd for 2(C₁₂H₁₇N₅O₄)•Na⁺ 613.2, found 613.1.

((2*S*,3*S*,5*R*)-3-azido-5-(2-ethoxy-5-methyl-4-oxopyrimidin-1(4*H*)-yl)tetrahydrofuran-2-yl)methyl acetate: Compound 4.



To a stirred solution of **3** (7.0 g, 23.7 mmol, 1 equiv) in acetic anhydride (200 mL) was added 4-dimethylaminopyridine (DMAP) (57 mg, 0.5 mmol, 0.02 equiv) at room temperature. The reaction mixture was stirred for 12 h, then the bulk of the solvent was removed by successive rounds of azeotropic co-evaporation with toluene. The resulting oil was re-dissolved in CH_2Cl_2 (500 mL), then washed with water (250 mL) and saturated aqueous NaCl solution (250 mL). The extracted organic layer was dried over Na₂SO₄ and concentrated under reduced pressure. The crude material was then purified by combiflash silica gel chromatography with a CH_2Cl_2 -MeOH elution mixture (0-to-10% MeOH gradient) to afford the 5'-*O*-acetylated nucleoside **4** (7.5 g, 95% yield).

¹**H NMR** (400 MHz, CDCl₃) δ 7.34 (q, J = 1.5 Hz, 1H), 6.12 (t, J = 6.5 Hz, 1H), 4.53 (q, J = 7.0 Hz, 2H), 4.40 (dd, J = 12.5, 4.0 Hz, 1H), 4.36 (dd, J = 12.5, 4.0 Hz, 1H), 4.18 (ddd, J = 7.5, 5.5, 5.0 Hz, 1H), 4.13 (ddd, J = 5.5, 4.0, 4.0 Hz, 1H), 2.51 (ddd, J = 14.0, 6.5, 5.0 Hz, 1H), 2.33 (ddd, J = 14.0, 7.5, 6.5 Hz, 1H), 2.16 (s, 3H), 2.00 (d, J = 1.5 Hz, 3H); and 1.41 (t, J = 7.0 Hz, 3H); ¹³**C NMR** (100 MHz, CD₃OD) δ 171.7, 170.3, 154.8, 131.9, 118.0, 85.6, 82.1, 65.4, 63.2, 60.4, 38.6, 20.9, 14.4, and 14.2; **LRMS** (ESI): calcd for C₁₄H₁₉N₅O₅•Na⁺ 360.128, found 360.13; calcd for 2(C₁₄H₁₉N₅O₅)•Na⁺ 697.27, found 697.31.

((2*S*,3*S*,5*R*)-3-amino-5-(5-methyl-4-oxo-2-thioxo-3,4-dihydropyrimidin-1(2*H*)-yl)tetrahydrofuran-2-yl)methyl acetate: Compound 6.



A representative reaction setup is depicted in Fig. S1 below. A two-neck reaction flask containing a solution of 4 (1.0 g, 3.0 mmol, 1 equiv) in anhydrous pyridine (30 mL) was purged with Ar (or N₂) for 30 min at room temperature. To this solution was added freshly distilled TMG (1.5 mL, 11.7 mmoles, 4 equiv), and the reaction mixture was cooled in an ice-bath. H₂S gas was then gently bubbled through a glass pipette into the stirred reaction solution, which turned to a deep green color over 5-10 mins. After 1 h, the H₂S gas source was closed, the remaining H₂S in the system was bubbled out, and the glass pipette was raised above the reaction liquid. After the pressure was equalized, valves connecting the tygon tubes to the reaction flask were closed, and

the reaction mixture was stirred at room temperature for 16 hrs. At the end of this mixing period, Ar (or N₂) was bubbled into the mixture for ca. 1 h to remove any residual volatile sulfur species. The reaction mixture was poured into a flask containing chloroform (300 mL). While being stirred vigorously, 1N HCl was added dropwise until the pH decreased to around 4. The chloroform layer was removed, and the remaining aqueous layer was treated with fresh chloroform. The stirred emulsion was neutralized by addition of saturated aqueous NaHCO₃ solution, and the organic part was then extracted and washed with water and saturated aqueous NaCl solution. The combined organic layers were dried over Na₂SO₄ and concentrated under reduced pressure. The resulting crude material was purified by combiflash silica gel chromatography with a CH₂Cl₂-MeOH elution mixture (0-to-15% MeOH gradient) to afford the 2-thio modified nucleoside **6** (470 mg, 53% yield).



Figure S1. Experimental Setup for the Sulfuration Reaction. Safety flask 1 prevents the reaction mixture from reaching into the H₂S source in case of an unexpected downstream pressure increase. Similarly, safety flask 2 prevents bleach from mixing into the reaction mixture in case of an unexpected pressure buildup.

Intermediate 5: ¹H NMR (400 MHz, CD₃OD) δ 7.58 (q, *J* = 1.0 Hz, 1H), 6.17 (dd, *J* = 6.5, 4.5 Hz, 1H), 4.49⁺ (dq, *J* = 2.5, 7.0 Hz, 1H), 4.49⁻ (dq, *J* = 2.5, 7.0 Hz, 1H), 4.43 (dd, *J* = 12.5, 5.0 Hz, 1H), 4.35 (dd, *J* = 12.5, 7.5 Hz, 1H), 3.95 (ddd, *J* = 7.5, 7.5, 5.0 Hz, 1H), 3.47 (q, *J* = 7.5 Hz, 1H), 2.37 (ddd, *J* = 14.0, 7.5, 4.5 Hz, 1H), 2.31 (ddd, *J* = 14.0, 7.5, 6.5 Hz, 1H), 2.10 (s, 3H), 1.97 (d, *J* = 1.0 Hz, 3H), and 1.42 (t, *J* = 7.0 Hz, 3H); ¹³C NMR (100 MHz, CD₃OD) δ 175.1, 172.5, 156.8,

136.3, 117.5, 88.0, 86.5, 66.7, 65.0, 52.0, 42.1, 20.9, 14.6, and 13.9; **LRMS** (ESI): calcd for $C_{14}H_{21}N_3O_5 \cdot H^+$ 312.16, found 311.95.

Product 6: ¹**H NMR** (400 MHz, CD₃OD) δ 7.74 (q, *J* = 1.5 Hz, 1H), 6.78 (dd, *J* = 7.0, 4.0 Hz, 1H), 4.51 (dd, *J* = 12.5, 5.0 Hz, 1H), 4.35 (dd, *J* = 12.5, 3.0 Hz, 1H), 3.95 (ddd, *J* = 7.5, 5.0, 3.0 Hz, 1H), 3.41 (q, *J* = 7.5 Hz, 1H), 2.40 (ddd, *J* = 14.0, 7.5, 7.0 Hz, 1H), 2.30 (ddd, *J* = 14.0, 7.5, 4.0 Hz, 1H), 2.13 (s, 3H), and 1.95 (d, *J* = 1.5 Hz, 3H); ¹³**C NMR** (100 MHz, CD₃OD) δ 176.0, 172.5, 163.4, 138.0, 116.6, 90.7, 86.6, 64.4, 51.8, 42.1, 21.0, and 13.0.; **LRMS** (ESI): calcd for C₁₂H₁₇N₃O₄S•H⁺ 300.10, found 299.53; calcd for C₁₂H₁₇N₃O₄S•H⁺ 322.08, found 321.75.

1-((2*R*,4*S*,5*S*)-4-amino-5-(hydroxymethyl)tetrahydrofuran-2-yl)-5-methyl-2-thioxo-2,3-dihydropyrimidin-4(1*H*)-one: Compound S1.



An NH₃ solution (7 N) in MeOH (8 mL) was introduced into a pressure reaction vessel containing a solution of **6** (450 mg, 1.5 mmol, 1 equiv) in MeOH (8 mL) at room temperature. The reaction solution was then stirred for 16 hours, and the solvent was removed by rotary evaporation and the residue was then left under high-vacuum for 2 hours. Dried deacetylated nucleoside **S1** (350 mg, 92% yield) was sufficiently pure (by ¹H NMR analysis) and was used directly in the next step without any purification.

¹**H NMR** (400 MHz, CD₃OD) δ 8.25 (q, *J* = 1.5 Hz, 1H), 6.77 (dd, *J* = 6.5, 4.0 Hz, 1H), 3.94 (dd, *J* = 12.5, 3.0 Hz, 1H), 3.83 (dd, *J* = 12.5, 3.0 Hz, 1H), 3.75 (ddd, *J* = 7.5, 3.0, 3.0 Hz, 1H), 3.51 (q, *J* = 7.5 Hz, 1H), 3.35 (s, 1H), 2.35 (ddd, *J* = 14.0, 7.5, 6.5 Hz, 1H), 2.30 (ddd, *J* = 14.0, 7.5, 4.0 Hz, 1H), and 1.92 (d, *J* = 1.5 Hz, 3H); ¹³**C NMR** (100 MHz, CD₃OD) δ 175.8, 163.6, 139.1, 116.2, 90.7, 89.5, 61.2, 50.5, 42.7, and 12.9; **LRMS** (ESI): calcd for C₁₀H₁₅N₃O₃S•H⁺ 258.09, found 257.86; calcd for [C₁₀H₁₅N₃O₃S - (H⁺)]⁻ 256.07, found 255.97.

1-((2*R*,4*S*,5*S*)-4-amino-5-(hydroxymethyl)tetrahydrofuran-2-yl)-5-methyl-2-thioxo-2,3-dihydropyrimidin-4(1*H*)-one: Compound 7.



To a stirred solution of **S1** (300 mg, 1.2 mmoles, 1 equiv) and 1 M aqueous Na₂CO₃ (0.7 mL, 0.6 equiv) in pyridine (12 mL) was added 9-fluorenylmethyl *N*-succinimidyl carbonate (Fmoc-OSu) (530 mg, 1.6 mmoles, 1.3 equiv) at 0 °C. The reaction mixture was stirred at room temperature for 6 h and then filtered through a PTFE filter (pore size 0.22 μ m). The filtrate was concentrated to remove the bulk pyridine, and the residue was then dissolved with CHCl₃, and then washed with a minimum amount of water. The organic layer was concentrated under reduced pressure to give an oil, which was triturated with the HPLC-grade acetonitrile. The resulting suspension was centrifuged and the supernatant was removed to provide the 3'-Fmoc protected nucleoside 7 (440 mg, 78% yield).

¹**H NMR** [400 MHz, CD₃S(O)CD₃] δ 12.2 (br s, 1H), 8.09 (s, 1H), 7.89 (d, J = 7.5 Hz, 2H), 7.69 (d, J = 7.5 Hz, 2H), 7.41 (t, J = 7.5 Hz, 2H), 7.33 (t, J = 7.5 Hz, 2H), 6.78 (br t, J = 6.0 Hz, 1H), 5.25 (br s, 1H), 4.35 (d, J = 6.5 Hz, 2H), 4.23 (t, J = 6.5 Hz, 1H), 4.15 (br ddd, J = 7.5, 6.5, 6.5 Hz, 1H), 3.88 (br m, 1H), 3.75–3.69 (nfom, 1H), 3.61–3.54 (nfom, 1H), 2.35–2.26 (m, 1H), 2.23–2.16 (m, 1H), and 1.82 (s, 3H); ¹**H NMR** [400 MHz, CD₃OD] δ 12.6 (br s, 1H), 8.11 (s, 1H), 7.89 (d, J = 7.5 Hz, 2H), 7.69 (d, J = 7.5 Hz, 2H), 7.41 (t, J = 7.5 Hz, 2H), 7.33 (t, J = 7.5 Hz, 2H), 6.77 (br t, J = 6.0 Hz, 1H), 5.25 (br t, J = 5.0 Hz, 1H), 4.35 (d, J = 6.5 Hz, 2H), 4.23 (t, J = 6.5 Hz, 1H), 4.15 (br ddd, J = 7.5, 6.5, 6.5 Hz, 1H), 3.88 (br m, 1H), 3.75–3.69 (br m, 1H), 3.61–3.54 (br m, 1H), 2.35–2.26 (m, 1H), 2.23–2.16 (m, 1H), and 1.83 (s, 3H); ¹³C **NMR** [100 MHz, CD₃S(O)CD₃] δ 174.1, 160.6, 156.0, 143.8 (2C), 140.7 (2C), 136.7, 127.6 (2C), 127.0 (2C), 125.1 (2C), 120.1 (2C), 115.1, 88.5, 85.1, 65.4, 60.1, 49.4, 46.7, 38.0, and 12.5; **LRMS** (ESI): calcd for C₂₅H₂₅N₃O₅S•Na⁺ 502.14, found 501.98; calcd for C₂₅H₂₅N₃O₅S•Cl⁻ 514.12, found 514.10.

3'- amino-3'-deoxy-2-thio-thymidine-5'-phosphor-2-methylimidazolide: 3'-NH₂-2-MeImpdds²T.



To a stirred solution of 7 (400 mg, 0.8 mmol, 1 equiv) in $PO(OMe)_3$ (8 mL) was added freshly activated 3Å molecular sieves and the mixture was stirred under Ar for 30 min. To this mixture was then added freshly distilled 2,6-lutidine (200 µL, 1.6 mmol, 2 equiv) and POCl₃ (120 µL, 1.2 mmol, 1.5 equiv) sequentially at 0 °C. The resulting white suspension was stirred for 2 hours and treated with 2-methylimidazole (260 mg, 3.2 mmol, 4 equiv). Resulting reaction mixture was allowed to warm to room temperature and stirred for an additional 2 hours. The mixture was then added into a pre-chilled mixture of acetone/diethyl ether/triethylamine/saturated sodium perchlorate solution in acetone (400:200:20:1 mL) in order to precipitate the desired product as a sodium salt. Most of the supernatant was removed by pipette-suction, and the remaining suspension was transferred into two falcon tubes, which were centrifuged to pellet the solid material. The supernatant was removed and a mixture of acetone/diethyl ether (1:1) was added into the tubes. After another centrifugation, the isolated solid material was purified by combiflash column (C18Aq) chromatography with an acetonitrile/TEAB buffer (pH ca. 7.5) elution mixture. The aqueous product fraction, which was determined by LR-ESI analysis, was placed into a falcon tube and freeze-dried under high-vacuum at -15 °C to afford the triethylammonium salt of the Fmoc protected 3'-amino-5'-phosphoroimidazolide intermediate. ³¹P NMR [161 MHz, CD₃OD/CD₃S(O)CD₃]: δ –9.2; LRMS (ESI): calcd for C₂₉H₃₀N₅O₇PS•Et₃NH⁺ 725.29, found 725.3; also calcd for $[C_{29}H_{30}N_5O_7PS - (H^+)]^-$ 622.15, found 622.1. To a stirred solution of this intermediate (100 mg, 0.13 mmol, 1 equiv) in DMF (2 mL), was added piperidine (130 µL, 1.3 mmol, 10 equiv) at 0 °C. The reaction mixture was stirred for 30 min at the same temperature, then directly concentrated in an ice-bath under high-vacuum. The resulting oil was immediately treated with a chilled mixture (21 mL) of acetone/diethyl ether/saturated sodium perchlorate solution in acetone (20:20:1 volume ratio) in order to precipitate the desired product. Most of the supernatant was removed by pipette-suction, and the remaining suspension was transferred into two falcon tubes, which were vortexed and then centrifuged to pellet the solid. The supernatant was removed

again by suction and a mixture of acetone/diethyl ether (1:1) was added into the tubes. After another vortexing and centrifugation, the liquid was removed and the solid material was purified by reverse-phase HPLC with an acetonitrile/TEAB buffer (pH ca. 7.5) elution mixture. The product fraction was transferred into a falcon tube and freeze-dried under high-vacuum at -15 °C to afford **3'-NH₂-2-MeImpdds²T** (43 mg, 26% overall yield from nucleoside **7**). The ¹H and ³¹P NMR spectroscopic data matched the previously reported results.² LRMS (ESI): calcd for [C₁₄H₂₀N₅O₅PS – (H⁺)]⁻ 400.09, found 399.96.



Figure S2. UV-VIS Spectrum of 3'-NH₂-2-MeImpdds²T































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Supporting Information













4. Monomer Decay-³¹P NMR Experiment

To determine the rate of decomposition of 3'-NH₂-2-MeImpdds²T under our primer extension reaction conditions, we have carried out a monomer decay-³¹P NMR experiment (Fig. S3).



Figure S3. Monomer decay-³¹P NMR spectral overlay of 3'-NH₂-2-MeImpdds²T. This nucleotide (10 mM) was dissolved in a solution (pH 7.5) of NaCl (150 mM), HEI (100 mM), CAPS:MES:HEPES (100 mM each) prepared using H₂O/D₂O (4:1). Each spectrum was acquired at 4 °C and referenced externally — by means of a co-axial NMR tube — to Ph₃PO dissolved in CDCl₃ at 26 ppm. Chemical shifts of 3'-NH₂-2-MeImpdds²T and its 5',3'-cyclization product appear at -11.9 and 1.7 ppm, respectively.

Results revealed that the half-life of 3'-NH₂-2-MeImpdds²T is 77 min, or 1.3 h (Fig. S4), which is similar to that (1.2 h) we previously measured for the unmodified analog, 3'-NH₂-2-MeImpddT.³ This suggests that the 2-thio modification does not affect the rate of cyclization of activated T monomer.



Figure S4. Plot of the percent decay of 3'-NH₂-2-MeImpdds²T. $R^2 = 0.982$.

5. Variable pH-¹³C NMR Experiment

We have carried out a variable pH-¹³C NMR spectroscopy experiment (Fig. S5) on 3'-amino-2',3'-dideoxy-2-thio-thymidine (S1) to measure the p K_a of the protonated form of 3'-amine. We have monitored the resonance of C3' of S1, while gradually increasing the pH of the NMR sample.



Figure S5. Variable pH-¹³C NMR spectral overlay of 3'-amino-2',3'dideoxy-2-thio-thymidine (S1). This nucleoside (50 mM) was dissolved in a phosphate buffer (300 mM) prepared using H_2O/D_2O (4:1). Each spectrum was acquired at 100 MHz and referenced externally — by means of a co-axial NMR tube — to $C(O)(CD_3)_2$ at 30.0 ppm (middle peak of the septet). The C3' resonance is observed to shift downfield from 83.4 to 87.0 ppm as the pH increases from 5.23 to 9.21, whereas the C4' resonance shifts minimally, from 89.0 to 89.5 ppm.



We have plotted the C3' chemical shift values (δ) against pH (Fig. S6) and fit to equation 1,⁴ derived from the Henderson-Hasselbalch equation,

$$\delta = \delta_{acid} + \frac{\delta_{base} \, 10^{(pH-pKa)}}{1 + 10^{(pH-pKa)}} \tag{1}$$

where δ_{acid} and δ_{base} correspond to the extrapolated C3' chemical shift values at the low and high

extremes of pH, respectively. With an excellent correlation coefficient ($R^2 = 0.999$), we have determined the midpoint of pH value as 7.51, which corresponds to p K_a of the 3'-amino group.



Figure S6. Plot of the chemical shift for C3' resonance of 3'-amino-2',3'-dideoxy-2-thio-thymidine with respect to pH.

6. LC-HRMS Analysis of the DNA Primer Strand.





Exact mass (Neg. mode): 5165.937; measured accurate mass: 5165.925, error: -2.3 ppm.



Figure S7. LC trace of the DNA primer strand.

7. Kinetic Studies of Primer Extension Reactions

Primer and template oligonucleotide strands were re-annealed by a heat-cool cycle in a PCR instrument. Primer extension reactions on poly(A) RNA templates (Fig. S8), which were performed in a 4 °C cold room, were initiated by addition of the monomer solution (for a total monomer concentration of 10 or 30 mM). Each reaction aliquot was quenched by 3x volume of formamide, and the quenched samples were stored at -80 °C until the gel electrophoresis.



Monomer soln. pH 7.9 (\pm 0.1); Reaction soln. pH 7.5 (\pm 0.1); CAPS:MES:HEPES (100 mM each) NaCl (150 mM) and HEI (100 mM); primer / template (0.2 μ M:1.0 μ M)

Figure S8. Primer extension kinetics.

8. X-Ray Crystallography

Nucleoside **6** was crystallized from chloroform. A crystal was mounted on a diffractometer and data was collected at 100 K. The intensities of the reflections were collected by means of a Bruker APEX II DUO CCD diffractometer (Cu_{K.} radiation, λ =1.54178 Å), equipped with an Oxford Cryosystems nitrogen flow apparatus. The collection method involved 1.0° scans in ω at -30°, -55°, -80°, 30°, 55°, 80° and 115° in 2 θ . Data integration down to 0.84 Å resolution was carried out using SAINT V8.34 C with reflection spot size optimization.⁵ Absorption corrections were made with the program SADABS.⁵ The structure (Fig. S9) was solved by the Intrinsic Phasing methods and refined by least-squares methods against F^2 using SHELXT-2014⁶ and SHELXL-2014 with OLEX 2 interface.⁷ Non-hydrogen atoms were refined anisotropically, and hydrogen atoms were allowed to ride on the respective atoms. Crystal data as well as details of data collection and refinement are summarized in Table S1, and geometric parameters are shown in Table S2. The Ortep plots produced with SHELXL-2014 program, and the other drawings were produced with Accelrys DS Visualizer 2.0.⁸



Figure S9. Crystal structure of nucleoside 6. (Right) Perspective views showing 50% probability displacement.

Crystal data			
Chemical formula	$C_{12}H_{17}N_3O_4S$ •CHCl ₃		
M _r	418.71		
Crystal system, space group	Orthorhombic, $P2_12_12_1$		
Temperature (K)	100		
<i>a</i> , <i>b</i> , <i>c</i> (Å)	4.9957 (3), 17.7935 (9), 20.4746 (10)		
$V(\text{\AA}^3)$	1820.01 (17)		
Ζ	4		
Radiation type	Cu <i>K</i> α		
μ (mm ⁻¹)	5.85		
Crystal size (mm)	$0.12 \times 0.01 \times 0.01$		
Data collection	·		
Diffractometer	Bruker D8 goniometer with CCD area detector diffractometer		
Absorption correction	Multi-scan SADABS		
T_{\min}, T_{\max}	0.683, 0.815		
No. of measured, independent and observed $[I > 2\sigma(I)]$ reflections	15691, 3129, 2776		
R _{int}	0.091		
$(\sin \theta / \lambda)_{max} (\text{\AA}^{-1})$	0.595		
	·		
Refinement			
$R[F^2 > 2\sigma(F^2)], wR(F^2), S$	0.079, 0.214, 1.08		
No. of reflections	3129		
No. of parameters	233		
No. of restraints	6		
H-atom treatment	H-atom parameters constrained		
$\Delta \rho_{\text{max}}, \Delta \rho_{\text{min}} (e \text{ Å}^{-3})$	0.73, -0.43		
Absolute structure	Flack x determined using 930 quotients [(I+)-(I-)]/[(I+)+(I-)] ⁹		
Absolute structure parameter	0.03 (3)		

Table S1. Experimental details for the X-ray analysis.

C1—N1:	1.342 (12)	N1—C1—S1:	120.2 (7)
C1—N2:	1.365 (12)	N2—C1—S1:	124.2 (7)
C1—S1:	1.684 (10)	O2—C5—N2:	107.2 (7)
C2—O1:	1.239 (11)	O2—C5—C6:	108.0 (7)
C2—N1:	1.394 (13)	N2—C5—C6:	114.6 (8)
C2—C3:	1.443 (13)	N3—C7—C6:	112.2 (7)
C3—C4:	1.337 (13)	N3—C7—C8:	111.4 (8)
C3—C10:	1.503 (14)	O3—C9—C8:	109.3 (8)
C5—N2:	1.485 (11)	S1—C1—N2—C5:	0.2 (13)
C7—N3:	1.470 (11)	O2—C5—N2—C4:	64.6 (10)

Table S2. Geometric parameters (Å, °). Most relevant measurements are listed.

9. References

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