

Supporting Information

Optimization of an Unnatural Base Pair Towards Natural-Like Replication

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	Contents	Page
Scheme S1	d 5FM synthetic scheme	S4
Scheme S2	d 5NaM synthetic scheme	S4
	Synthetic procedures and characterizations	S5–S7
Table S1	Kinetic data for insertion of triphosphates opposite d 5SICS in sequence context I	S8
Table S2	Kinetic data for insertion of triphosphates opposite d MMO2 , d 5FM , or d NaM in sequence context I	S9
Table S3	Kinetic data for extension of dN:d 5SICS in sequence context I	S10
Table S4	Kinetic data for extension of dN:d MMO2 , dN:d 5FM , and dN:d NaM in sequence context I	S11
Table S5	Kinetic data for insertion of triphosphates opposite d 5SICS in sequence context II	S12
Table S6	Kinetic data for insertion of triphosphates opposite d MMO2 , d 5FM , or d NaM in sequence context II	S13
Table S7	Kinetic data for extension of dN:d 5SICS in sequence context II	S14
Table S8	Kinetic data for extension of dN:d MMO2 , dN:d 5FM , and dN:d NaM in sequence context II	S15
Table S9	Kinetic data for insertion of d NaM opposite d 5SICS in sequence context III	S16
Table S10	Kinetic data for insertion of d NaM opposite d 5SICS in sequence context IV	S16
Figure S1	Raw gel data and plots of representative data(d NaMTP insertion opposite d 5SICS and extension in sequence context I and II)	S17

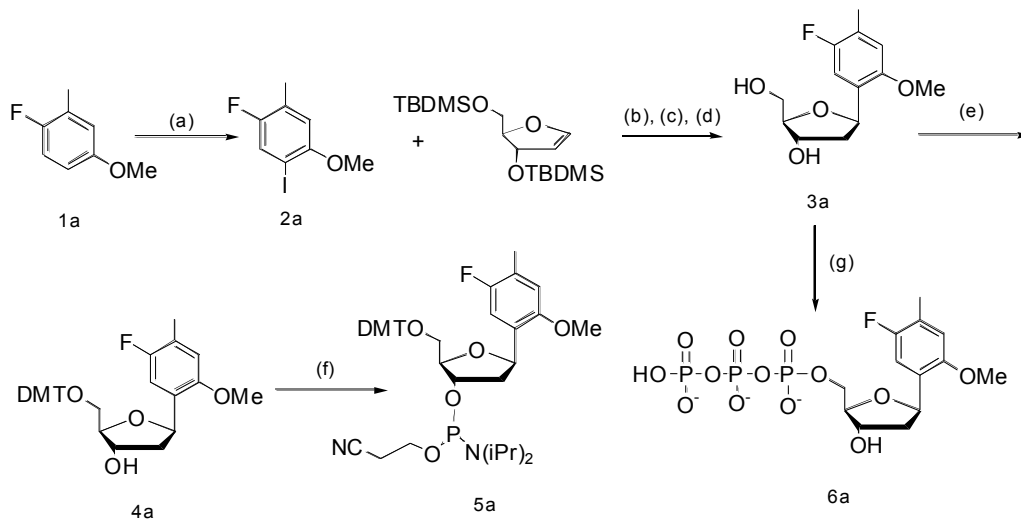
NMR and HRMS Spectrum of Novel Compounds

Compound (see scheme S1 and S2)	Spectrum	Page
3a	^1H NMR	S18
3a	C^{13} NMR	S19
3a	Homo-Cozy NMR	S20
3a	1D-NOE NMR	S21
4a	^1H NMR	S22
4a	C^{13} NMR	S23
5a	^1H NMR	S24
5a	P^{31} NMR	S25
6a	P^{31} NMR	S26
dNaM intermediate	^1H NMR	S27
dNaM intermediate	C^{13} NMR	S28
dNaM intermediate	Homo-Cozy NMR	S29
dNaM intermediate	1D-NOE NMR	S30
2b	^1H NMR	S31
2b	C^{13} NMR	S32
3b	^1H NMR	S33
3b	C^{13} NMR	S34
4b	^1H NMR	S35
4b	P^{31} NMR	S36
5b	P^{31} NMR	S37
3a	HRMS	S38
4a	HRMS	S39
5a	HRMS	S40
2b	HRMS	S41
3b	HRMS	S42
4b	HRMS	S43

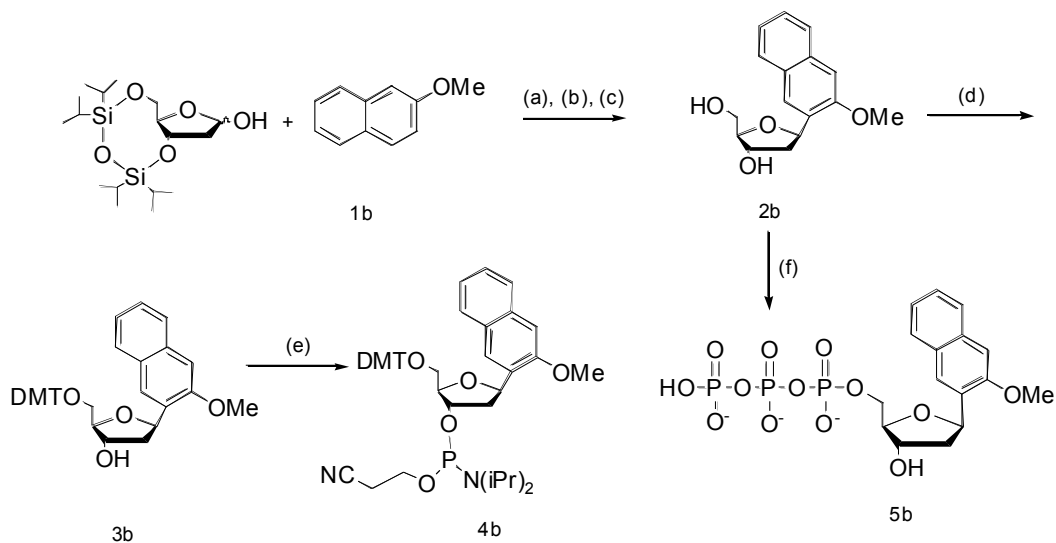
MALDI-TOF mass spectral data^a

Sequence	MALDI-TOF signal [M ⁺]		Page
	Calc. m/z	Found m/z	
P1: 5'-d (TAATACGACTCACTATAGGGAGANaM)	7416	7413	S44
P2: 5'-d (TAATACGACTCACTATAGGGAGAFM)	7398	7394	S45
P3: 5'-d (TAATACGACTCACTATAGGGAGA5SICS)	7433	7426	S46
P4: 5'-d (TAATACGACTCACTATAGGGAGAMMO2)	7380	7374	S47
T1: 3'-d (ATTATGCTGAGTGATATCCCTCTNaMGCTAGGTTCGGCAGGATCGC)	13890	13893	S48
T2: 3'-d (ATTATGCTGAGTGATATCCCTCTFMGCTAGGTTCGGCAGGATCGC)	13872	13876	S49
T3: 3'-d (ATTATGCTGAGTGATATCCCTCT5SICSGCTAGGTTCGGCAGGATCGC)	13907	13907	S50
T4: 3'-d (ATTATGCTGAGTGATATCCCTCTMMO2GCTAGGTTCGGCAGGATCGC)	13854	13857	S51
P5: 5'-d (TAATACGACTCACTATAGGGAGCNaM)	7393	7396	S52
P6: 5'-d (TAATACGACTCACTATAGGGAGCFM)	7375	7382	S53
P7: 5'-d (TAATACGACTCACTATAGGGAGC5SICS)	7410	7414	S54
P8: 5'-d (TAATACGACTCACTATAGGGAGCMMO2)	7357	7361	S55
T5: 3'-:d (ATTATGCTGAGTGATATCCCTCGNaMTCTAGGTTCGGCAGGATCGC)	13890	13891	S56
T6: 3'-:d (ATTATGCTGAGTGATATCCCTCGFMTCTAGGTTCGGCAGGATCGC)	13872	13877	S57
T7: 3'-:d (ATTATGCTGAGTGATATCCCTCG5SICSTCTAGGTTCGGCAGGATCGC)	13907	13888	S58
T8: 3'-:d (ATTATGCTGAGTGATATCCCTCGMMO2TCTAGGTTCGGCAGGATCGC)	13854	13852	S59
T9: 3'-d (ATTATGCTGAGTGATATCCCTCA5SICSTCTAGGTTCGGCAGGATCGC)	13916	13915	S60
T10: 3'-d (ATTATGCTGAGTGATATCCCTCC5SICSTCTAGGTTCGGCAGGATCGC)	13892	13890	S61

^a **P1-P4:** primers for context I, **T1-T4:** templates for context I, **P5-P8:** primers for context II, **T5-T8:** templates for context II and **T9-T10:** templates for context III and IV.



Scheme S1. (a) I_2 , $AgNO_3$, MeOH, yield:73%; (b) $Pd(OAc)_2$, $AsPh_3$, $n-Bu_3N$, DMF, 70 °C ; (c) n -TBAF, 0 °C; (d) $NaBH(OAc)_3$, AcOH, CH_3CN , overall yield:52%; (e) DMTrCl, DMAP, Et_3N , pyridine, yield:78%; (f) $NC(CH_2)_2OP(iPr_2)N_2$, DIPEA, CH_2Cl_2 , yield:85%; (g) Proton Sponge, $POCl_3$, Bu_3N , Bu_3NPPi , $(MeO)_3P$, DMF, -20 °C.



Scheme S2. (a) 1.6 M $n-BuLi$, THF, -10 °C; (b) $n-Bu_3P$, TMAD, THF, rt ; (c) TBAF, THF, rt, overall yield: 22%; (d) DMTrCl, DMAP, Et_3N , pyridine, yield: 82%; (e) $NC(CH_2)_2OP(iPr_2)N_2$, DIPEA, CH_2Cl_2 , yield: 92%; (f) Proton Sponge, $POCl_3$, Bu_3N , Bu_3NPPi , $(MeO)_3P$, DMF, -20 °C.

Synthetic procedures and characterizations

Compound 2b. A mixture of 2-methoxy naphthalene (700mg, 4.4 mmol) in THF (10 mL) was stirred under nitrogen atmosphere at room temperature for 30 minutes, after which 1.6 M *n*-BuLi (2.7ml, 28.5mmol) was added dropwise and the mixture was stirred for 2 h at room temperature. After cooling to -10 °C, 3,5-O-[1,1,3,3-tetrakis(1-methylethyl)-1,3-disiloxanediy] 2-deoxy-D-erythro-pent-1-enitol (537 mg, 1.43 mmol) was added and the temperature was increased slowly to room temperature and the mixture stirred for 30 min. The reaction mixture was extracted with EtOAc (50 mL × 3) and saturated NH₄Cl aq. (50 mL) and the combined organic layers were dried over anhydrous Na₂SO₄. After filtration and evaporation, the residue was purified by silica gel column chromatography (EtOAc/hexane (1:2), R_f = 0.38). The eluted product, without further purification, was dissolved in THF (3 mL) and tri-*n*-butyl phosphin (87 µl, 0.35 mmol) was added. The resulting solution was cooled to 0 °C, *N,N,N',N'*-tetramethylazodicarboxamide (60 mg, 0.35 mmol) was added, and the solution was stirred overnight at room temperature. Tetrabutylammoniumfluoride (0.45 mL, 0.45 mmol) was added to this mixture at room temperature and stirred for 3 h. The reaction mixture extracted with EtOAc (30 mL × 3) and saturated NaHCO₃ aq. (30 mL) and the combined organic layers were dried over anhydrous Na₂SO₄. After filtration and evaporation, the residue was purified by silica gel column chromatography (EtOAc R_f = 0.42) to afford **2b** (86 mg, 0.31 mmol, 22% from 3 steps) as a white solid. ¹H NMR (300 MHz, MeOH-*d*4) δ 7.28-8.07 (m, 6 H, ArH), 5.57 (dd, *J*=9.9 Hz, 5.7 Hz, 1 H, H-1'), 4.37-4.41 (m, 1 H, H-3'), 4.05-4.10 (m, 1 H, H-4'), 4.01 (s, 3H, OCH₃), 4.82 (dd, *J*=5.4 Hz, *J*=2.7 Hz, 2H, H-5'), 2.49-2.56 (m, 1 H, H-2'), 1.86-1.96 (m, 1 H, H-2'). ¹³C NMR (300 MHz, MeOH-*d*4) δ 155.1, 134.1, 129.0, 127.4, 126.2, 125.7, 124.5, 123.4, 104.7, 87.4, 75.7, 73.1, 62.9, 54.5, 48.6, 42.2. HRMS calcd for C₁₆H₁₈O₄ (MNa⁺) 297.1097, found 297.1104.

Compound 3a. A mixture of palladium acetate (16 mg, 0.071 mmol) and triphenylarsine (43.6 mg, 0.14 mmol) in DMF (3 mL) was stirred under argon atmosphere at room temperature for 20 min. Then **2a** (190 mg, 0.71 mmol), 1,4-anhydro-3,5-bis-O-(tert-butyl)dimethylsilyl)-2-deoxy-D-erythro-pent-1-enitol (205 mg, 0.594 mmol), and tri-*n*-butylamine (170mg, 0.92 mmol) in DMF (5 mL) were added, and the resulting reaction mixture was stirred under nitrogen at 70 °C for 18 h. The mixture was cooled to 0 °C, and 1 M tetra-*n*-butylammonium fluoride (3.16mL, 3.16 mmol) in THF was added and the mixture stirred for 1 h. The reaction mixture was filtered through Celite and extracted with EtOAc (50 mL × 3) and saturated NaHCO₃ aq. (50 mL). The combined organic layers were dried over anhydrous Na₂SO₄. After filtration and evaporation, the residue was purified by silica gel column chromatography (5% MeOH/CH₂Cl₂, R_f = 0.42). The eluted product was dissolved in acetic acid (2.5 mL) and acetonitril (2.5 mL). The solution was cooled to 0 °C, and sodium triacetoxyhydride (240 mg, 1.13 mmol) was added followed by stirring for 1 h. The reaction mixture was extracted with EtOAc (50 mL × 3) and saturated NaHCO₃ aq. (50 mL) and the combined organic layers were dried over

anhydrous Na₂SO₄. After filtration and evaporation, the residue was purified by silica gel column chromatography (EtOAc, R_f = 0.53) to afford **3a** (79 mg, 0.3 mmol, 52 % over 3 steps) as a white solid. ¹H NMR (300 MHz, MeOH-*d*₄) δ 7.26 (d, *J*=10.5 Hz, 1 H, ArH), 6.83 (d, *J*=6 Hz, 1 H ArH), 5.41 (dd, *J*=10.2 Hz, 5.7 Hz, 1 H, H-1'), 4.34 (dt, *J*=6 Hz, *J*=2.7 Hz 1 H, H-3'), 3.98(td, 1 H, *J*=5.4 Hz, *J*=2.7 Hz 1 H, H-4'), 3.86(s, 3H, OCH₃), 3.74 (dd, *J*=5.4 Hz, *J*=1.2 Hz, 2H, H-5'), 2.34-2.41 (m, 1H, H-2'), 2.30 (d, *J*=2.1 Hz, CH₃), 1.75-1.85(m, 1H, H-2'). ¹³C NMR (300 MHz, MeOH-*d*₄) δ 157.3, 130.1, 123.5, 112.9, 112.4, 87.4, 74.8, 73.2, 62.8, 55.2, 47.3, 42.2, 13.4. HRMS calcd for C₁₃H₁₇FO₄ (MNa⁺) 279.1002, found 279.1003.

General DMTr-protection Procedure. To a solution of free nucleoside (~100 mg, 1 eq.) and *N,N*-dimethylaminopyridine (0.2 eq.) in pyridine (1 mL), 4,4'- dimethoxytrityl chloride (1.3 eq.) and triethylamine (1.3 eq.) were added. The solution was stirred for 18 h at ambient temperature. After quenching the reaction by adding MeOH, the mixture was evaporated to dryness *in vacuo*. The residue was extracted with EtOAc (50 mL x 3) and water (50 mL), and the organic layer was collected, dried over anhydrous Na₂SO₄, filtered, and evaporated to dryness under reduced pressure. The crude product was purified by silica gel column chromatography (hexane:EtOAc = 3:1 to 2:1) to afford the DMTr-protected nucleoside as a white foam.

Compound 4a. ¹H NMR (300 MHz, CDCl₃) δ 7.12-7.41 (m, 11H, Ar-H), 6.81-6.89 (m, 4 H, Ar-H), 6.61(bs, 1 H, OH), 5.32 (dd, *J*=9.9 Hz, 5.7 Hz, 1 H, H-1'), 4.29-4.33 (m, 1 H, H-3'), 4.10-4.18(m, 1 H, H-4'), 3.83(bs, 6H, Ar-OCH₃), 3.71 (bs, 3H, Ar-OCH₃), 3.35(dd, 1H, *J*=9.6 Hz, *J*=4.8 Hz, H-5'), 3.20(dd, 1H, *J*=9.6 Hz, *J*=5.4 Hz, H-5'), 2.28-2.35 (m, 1H, H-2'), 2.21(s, 3H, CH₃) 1.79-1.88(m, 1H, H-2'). ¹³C NMR (300 MHz, CDCl₃) δ 158.6, 155.1, 151.1, 145.2, 135.4, 130.2, 129.5, 128.2, 127.5, 125.3, 112.3, 111.2, 87.4, 78.7, 76.9, 76.3, 73.1, 62.8, 55.8, 42.1, 14.5. HRMS calcd for C₃₄H₃₅FO₆ (MNa⁺) 581.2310, found 581.2312.

Compound 3b. ¹H NMR (300 MHz, CDCl₃) δ 7.09-7.64 (m, 11 H, Ar-H), 6.72-6.75 (m, 4 H, Ar-H), 5.44 (dd, *J*=9.0 Hz, 6.3 Hz, 1 H, H-1'), 4.30-4.34 (m, 1 H, H-3'), 3.97-4.07(m, 1 H, H-4'), 3.83(s, 6H, Ar-OCH₃), 3.68 (s, 3H, Ar-OCH₃), 3.36(dd, 1H, *J*=9.9 Hz, *J*=4.5 Hz, H-5'), 3.26(dd, 1H, *J*=9.6 Hz, *J*=5.4 Hz, H-5'), 2.36-2.43 (m, 1H, H-2'), 1.83-1.96(m, 1H, H-2'). ¹³C NMR (300 MHz, CDCl₃) δ 158.7, 155.0, 145.0, 136.2, 133.7, 132.7, 130.7, 128.3, 127.9, 126.9, 124.8, 123.7, 113.2, 104.9, 86.3, 85.6, 77.5, 75.5, 74.7, 64.5, 55.4, 42.4. HRMS calcd for C₃₇H₃₆O₆ (MNa⁺) 599.2404, found 599.2402.

General Phosphoramidite Synthesis Procedure. To a solution of DMTr-protected nucleoside (100 mg, 1 eq.) and diisopropylethylamine (4 eq.) in dichloromethane (1 mL) was added 2-cyanoethyl *N,N*-diisopropylchlorophosphoramidite (1.5 eq.) at 0 °C. After stirring for 30 min at ambient temperature, the reaction mixture was extracted with EtOAc (30 mL x 3) and saturated NaHCO₃ aq. (20 mL). The organic layers were collected, dried over anhydrous Na₂SO₄, filtered, and evaporated to dryness under reduced pressure. The

crude product was purified by silica gel column chromatography (hexane:EtOAc = 3:1 to 2:1) to afford compound as a pale yellow foam.

Compound 5a. ^1H NMR (400 MHz, CDCl_3) δ 7.18-7.47 (m, 11 H, Ar-H), 6.80-6.84 (m, 4 H, Ar-H), 6.5-6.6 (m, 1 H, Ar-H), 5.32(ddd, $J=7.8$ Hz, 5.7 Hz, 3.6Hz, 1 H, H-1'), 4.36-4.49 (m, 1 H, H-3'), 4.17-4.20(m, 1 H, H-4'), 3.76(t, 6H, Ar-OCH₃), 3.74 (bs, 3H, Ar-OCH₃), 3.64-3.83(m, 2H, CH₂), 3.50-3.62(m, 2H, CH₂), 3.22-3.27 (m, 2H, H-5'), 2.58-2.78 (m, 2H, CH), 2.41-2.52 (m, 1H, H-2'), 2.23(s, 3H, CH₃), 1.73-1.82(m, 1H, H-2'), 1.04-1.27 (m, 12 H, N-CH₃). ^{31}P NMR (400 MHz, CDCl_3): δ 148.5, 148.1. HRMS calcd for $\text{C}_{43}\text{H}_{52}\text{FN}_2\text{O}_7\text{P}(\text{MH}^+)$ 759.3569, found 759.3559.

Compound 4b. ^1H NMR (400 MHz, CDCl_3) δ 7.10-8.08 (m, 11 H, Ar-H), 6.80-6.83 (m, 4 H, Ar-H), 5.51-5.56 (m, 1 H, H-1'), 4.40-4.57 (m, 1 H, H-3'), 4.20-4.28(m, 1 H, H-4'), 3.93(d, 3H, Ar-OCH₃), 3.77 (bs, 6H, Ar-OCH₃), 3.65-3.73(m, 2H, CH₂), 3.54-3.66(m, 2H, CH₂), 3.30-3.43 (m, 2H, H-5'), 2.61-2.67(m, 2H, CH), 2.47-2.51(m, 1H, H-2'), 1.99(s, 3H, CH₃), 1.91-1.98(m, 1H, H-2'), 1.06-1.31 (m, 12 H, N-CH₃). ^{31}P NMR (400MHz, CDCl_3): δ 148.7, 148.1. HRMS calcd for $\text{C}_{46}\text{H}_{53}\text{N}_2\text{O}_7\text{P}(\text{MH}^+)$ 777.3663, found 777.3659.

General Procedure for Triphosphate Synthesis. Proton sponge (1.5 equiv) and the free nucleoside (1 equiv) were dissolved in trimethyl phosphate (0.3 M) and cooled to -20 °C. POCl_3 (1.5 equiv) was added dropwise, and the purple slurry was stirred at -20 °C for 2 h. Tributylamine (6.2 equiv) was added, followed by a solution of tributylammonium pyrophosphate (5.0 equiv) in DMF (0.5 M). After 5 min, the reaction was quenched by addition of 0.5 M aqueous $\text{Et}_3\text{NH}_2\text{CO}_3$ (20 vol-equiv) and the resulting solution was lyophilized. Purification by reverse-phase (C18) HPLC (4-35% CH_3CN in 0.1 M $\text{Et}_3\text{NH}_2\text{CO}_3$, pH 7.5) followed by lyophilization afforded the triphosphate as a white solid.

Compound 6a. ^{31}P NMR (400 MHz, D_2O): δ -9.00 – -9.11(m; γ -P), -10.64 (d, $J = 37.2$ Hz; α -P), -22.5 (t, $J = 37.5$ Hz; β -P).

Compound 5b. ^{31}P NMR (400 MHz, D_2O): δ -10.02 – -10.83 (m; γ -P, α -P), -22.65 - -22.98 (m; β -P).

Table S1. Kinetic data for insertion of dYTP d**5SICS** in sequence context I^a

5' -d (TAATACGACTCACTATAGGGAGA)

3' -d (ATTATGCTGAGTGATATCCCTCT**X**GCTAGGTTACGGCAGGATCGC)

X(Template)	dYTP	k_{cat} (min ⁻¹)	K_M (μM)	k_{cat}/K_M (M ⁻¹ min ⁻¹)
5SICS	MMO2	12 \pm 1	33 \pm 6	3.6 \times 10 ⁵
	5FM	8.2 \pm 1.8	2.3 \pm 0.3	3.6 \times 10 ⁶
	NaM	25 \pm 5	4.9 \pm 1.2	5.0 \times 10 ⁶
	5SICS	1.7 \pm 0.5	63 \pm 4	2.7 \times 10 ⁴
	A	1.2 \pm 0.3	54 \pm 6	2.2 \times 10 ⁴
	C	nd ^b	nd ^b	<1.0 \times 10 ³
	G	4.6 \pm 0.6	36 \pm 4	1.3 \times 10 ⁵
T	1.7 \pm 0.2	130 \pm 19	1.3 \times 10 ⁴	

^a See Experimental Section for experimental details. ^b Rates too slow to independently determine k_{cat} and K_M .

Table S2. Kinetic data for insertion of dYTP opposite dMMO2, d5FM, or dNaM in sequence context I^a

5'-d (TAATACGACTCACTATAGGGAGA)

3'-d (ATTATGCTGAGTGATATCCCTCT**X**GCTAGGTTACGGCAGGATCGC)

X(Template)	dYTP	k_{cat} (min ⁻¹)	K_M (μM)	k_{cat}/K_M (M ⁻¹ min ⁻¹)
MMO2	5SICS	8.5 ± 0.1	0.18 ± 0.01	4.7 × 10 ⁷
	MMO2	5.1 ± 0.7	44 ± 4	1.2 × 10 ⁵
	A	3.3 ± 0.3	32 ± 4	1.0 × 10 ⁵
	G	nd ^b	nd ^b	<1.0 × 10 ³
	C	nd ^b	nd ^b	<1.0 × 10 ³
	T	nd ^b	nd ^b	<1.0 × 10 ³
5FM	5SICS	3.3 ± 1.0	0.23 ± 0.03	1.4 × 10 ⁷
	FM	10.8 ± 2.4	12.2 ± 3.9	8.9 × 10 ⁵
	A	6.1 ± 1.0	20 ± 2	3.0 × 10 ⁵
	G	nd ^b	nd ^b	<1.0 × 10 ³
	C	nd ^b	nd ^b	<1.0 × 10 ³
	T	0.31 ± 0.1	131 ± 15	2.4 × 10 ³
NaM	5SICS	3.7 ± 0.8	0.10 ± 0.03	3.7 × 10 ⁷
	NaM	6.9 ± 1.6	2.0 ± 0.4	3.4 × 10 ⁶
	A	8.1 ± 1.7	11.4 ± 4.3	7.1 × 10 ⁵
	G	nd ^b	nd ^b	<1.0 × 10 ³
	C	nd ^b	nd ^b	<1.0 × 10 ³
	T	0.77 ± 0.12	136 ± 4.3	5.6 × 10 ³

^a See Experimental Section for experimental details. ^b Rates too slow to independently determine k_{cat} and K_M .

Table S3. Kinetic data for extension of dN:d**5SICS** by dCTP insertion in sequence context I^a

5' -d (TAATACGACTCACTATAGGGAGAY)

3' -d (ATTATGCTGAGTGATATCCCTCT**X**GCTAGGTTACGGCAGGATCGC)

X(Template)	Y	k_{cat} (min ⁻¹)	K_M (μM)	k_{cat}/K_M (M ⁻¹ min ⁻¹)
5SICS	MMO2	6.4 ± 1.1	3.4 ± 0.3	1.9 × 10 ⁶
	5FM	3.49 ± 0.11	0.63 ± 0.10	5.5 × 10 ⁶
	NaM	3.1 ± 0.9	2.47 ± 0.40	1.2 × 10 ⁶
	5SICS	nd ^b	nd ^b	<1.0 × 10 ³
	A	0.79 ± 0.01	76 ± 11	1.0 × 10 ⁴
	G	0.62 ± 0.02	127 ± 8	4.9 × 10 ³
	C	0.18 ± 0.01	44 ± 9	4.2 ± 10 ³
	T	2.0 ± 0.1	4.9 ± 0.3	4.0 ± 10 ⁵

^a See Experimental Section for experimental details. ^b Rates too slow to independently determine k_{cat} and K_M .

Table S4. Kinetic data for extension of dN:d**MMO2**, dN:d**5FM**, and dN:d**NaM** by dCTP insertion in sequence context I^a

5' -d (TAATACGACTCACTATAGGGAGAY)

3' -d (ATTATGCTGAGTGATATCCCTCT**X**GCTAGGTTACGGCAGGATCGC)

X(Template)	Y	k_{cat} (min ⁻¹)	K_M (μM)	k_{cat}/K_M (M ⁻¹ min ⁻¹)
MMO2	5SICS	3.8 ± 0.3	5.7 ± 1.2	6.7 × 10 ⁵
	MMO2	0.87 ± 0.28	165 ± 47	5.3 × 10 ³
	A	8.7 ± 0.9	187 ± 17	4.6 × 10 ⁴
	G	nd ^b	nd ^b	<1.0 × 10 ³
	C	3.90 ± 1.32	3.17 ± 0.23	1.2 × 10 ⁶
	T	6.23 ± 1.01	9.45 ± 1.23	6.6 × 10 ⁵
FM	5SICS	3.2 ± 0.7	1.4 ± 0.3	2.3 × 10 ⁶
	5FM	2.57 ± 0.39	99.5 ± 17.9	2.6 × 10 ⁴
	A	5.49 ± 0.08	17.3 ± 1.6	3.2 × 10 ⁵
	G	nd ^b	nd ^b	<1.0 × 10 ³
	C	2.4 ± 0.23	1.19 ± 0.10	2.0 × 10 ⁶
	T	4.14 ± 0.19	2.06 ± 0.17	2.0 × 10 ⁶
NaM	5SICS	5.73 ± 0.51	2.15 ± 0.50	2.7 × 10 ⁶
	NaM	0.78 ± 0.21	69 ± 25	1.1 × 10 ⁴
	A	3.99 ± 1.46	102 ± 35	3.9 × 10 ⁴
	G	nd ^b	nd ^b	<1.0 × 10 ³
	C	7.29 ± 0.69	4.78 ± 0.24	1.5 × 10 ⁶
	T	4.73 ± 0.42	8.33 ± 0.27	5.7 × 10 ⁵

^a See Experimental Section for experimental details. ^b Rates too slow to independently determine k_{cat} and K_M .

Table S5. Kinetic data for insertion of dYTP opposite d**5SICS** in sequence context II^a

5' -d (TAATACGACTCACTATAGGGAGC)

3' -d (ATTATGCTGAGTGATATCCCTCG**X**TCTAGGTTACGGCAGGATCGC)

X (Template)	dYTP	k_{cat} (min ⁻¹)	K_M (μM)	k_{cat}/K_M (M ⁻¹ min ⁻¹)
5SICS	MMO2	2.91 \pm 0.76	6.04 \pm 1.9	4.8 \times 10 ⁵
	5FM	9.52 \pm 1.14	2.29 \pm 0.21	4.2 \times 10 ⁶
	NaM	15.0 \pm 1.10	0.37 \pm 0.07	4.1 \times 10 ⁷
	5SICS	1.41 \pm 0.04	5.84 \pm 0.20	2.4 \times 10 ⁵
	A	1.23 \pm 0.24	20.8 \pm 2.86	5.9 \times 10 ⁴
	C	nd ^b	nd ^b	<1.0 \times 10 ³
	G	3.11 \pm 0.53	15.9 \pm 2.34	1.9 \times 10 ⁵
T	0.62 \pm 0.20	74.3 \pm 42.4	8.5 \times 10 ³	

^a See Experimental Section for experimental details. ^b Rates too slow to independently determine k_{cat} and K_M .

Table S6. Kinetic data for insertion of dYTP opposite dMMO2, d5FM, or dNaM in sequence context II^a

5' -d (TAATACGACTCACTATAGGGAGC)

3' -d (ATTATGCTGAGTGATATCCCTCG**X**TCTAGGTTACGGCAGGATCGC)

X(Template)	dYTP	k_{cat} (min ⁻¹)	K_M (μ M)	k_{cat}/K_M (M ⁻¹ min ⁻¹)
MMO2	5SICS	10.7 \pm 2.4	0.16 \pm 0.04	6.6 \times 10 ⁷
	MMO2	15.2 \pm 3.4	8.34 \pm 3.8	1.8 \times 10 ⁶
	A	17.4 \pm 4.1	10.2 \pm 3.3	1.7 \times 10 ⁶
	G	0.16 \pm 0.001	20.3 \pm 5.0	7.9 \times 10 ³
	C	0.65 \pm 0.04	219 \pm 46	3.0 \times 10 ³
	T	1.21 \pm 0.06	235 \pm 123	5.2 \times 10 ³
5FM	5SICS	12.3 \pm 2.6	0.15 \pm 0.04	8.0 \times 10 ⁷
	5FM	5.37 \pm 0.33	1.98 \pm 0.39	2.7 \times 10 ⁶
	A	22.6 \pm 4.1	7.4 \pm 1.5	3.1 \times 10 ⁶
	G	0.16 \pm 0.02	135 \pm 52	1.2 \times 10 ³
	C	1.15 \pm 0.01	152 \pm 88	7.5 \times 10 ³
	T	2.95 \pm 0.64	253 \pm 158	1.2 \times 10 ⁴
NaM	5SICS	12.4 \pm 3.6	0.22 \pm 0.04	5.5 \times 10 ⁷
	NaM	5.36 \pm 0.99	1.11 \pm 0.08	4.8 \times 10 ⁶
	A	21.9 \pm 4.2	15.3 \pm 1.2	1.4 \times 10 ⁶
	G	0.1 \pm 0.003	26.68 \pm 3.2	3.7 \times 10 ³
	C	0.63 \pm 0.19	98.3 \pm 10.4	6.4 \times 10 ³
	T	1.18 \pm 0.35	141 \pm 17.9	8.3 \times 10 ³

^a See Experimental Section for experimental details. ^b Rates too slow to independently determine k_{cat} and K_M .

Table S7. Kinetic data for extension of dN:d**5SICS** by dATP insertion in sequence context II^a

5'-d (TAATACGACTCACTATAGGGAGC**Y**)

3'-d (ATTATGCTGAGTGATATCCCTCG**X**TCTAGGTTACGGCAGGATCGC)

X(Template)	Y	k_{cat} (min^{-1})	K_M (μM)	k_{cat}/K_M ($\text{M}^{-1} \text{min}^{-1}$)
5SICS	MMO2	4.33 ± 1.17	13.5 ± 3.79	3.2×10^5
	5FM	7.28 ± 1.41	5.82 ± 1.70	1.3×10^6
	NaM	2.15 ± 0.18	3.15 ± 1.29	6.8×10^5
	5SICS	nd ^b	nd ^b	$<1.0 \times 10^3$
	A	0.10 ± 0.03	29.4 ± 15.9	3.6×10^3
	G	0.212 ± 0.04	324 ± 35	6.5×10^2
	C	nd ^b	nd ^b	$<1.0 \times 10^3$
	T	4.99 ± 0.32	12.0 ± 0.3	3.3×10^5

^a See Experimental Section for experimental details. ^b Rates too slow to independently determine k_{cat} and K_M .

Table S8. Kinetic data for extension of dN:d**MMO2**, dN:d**5FM**, and dN:d**NaM** by dATP insertion in sequence context II^a

5' -d (TAATACGACTCACTATAGGGAGC**Y**)

3' -d (ATTATGCTGAGTGATATCCCTCG**X**TCTAGGTTACGGCAGGATCGC)

X(Template)	Y	k_{cat} (min ⁻¹)	K_M (μM)	k_{cat}/K_M (M ⁻¹ min ⁻¹)
MMO2	5SICS	12.4 ± 1.44	7.30 ± 2.39	1.7 × 10 ⁶
	MMO2	nd ^b	nd ^b	<1.0 × 10 ³
	A	0.42 ± 0.12	38.4 ± 12.5	1.1 × 10 ⁴
	G	nd ^b	nd ^b	<1.0 × 10 ³
	C	4.91 ± 0.56	11.2 ± 2.41	4.4 × 10 ⁵
	T	6.67 ± 0.99	3.37 ± 0.51	2.0 × 10 ⁶
5FM	5SICS	13.2 ± 0.31	1.63 ± 0.02	8.2 × 10 ⁶
	5FM	nd ^b	nd ^b	<1.0 × 10 ³
	A	0.59 ± 0.06	13.4 ± 4.0	4.4 × 10 ⁴
	G	nd ^b	nd ^b	<1.0 × 10 ³
	C	7.17 ± 0.60	4.76 ± 0.77	1.5 × 10 ⁶
	T	6.97 ± 1.46	1.72 ± 0.23	4.1 × 10 ⁶
NaM	5SICS	17.3 ± 5.9	4.16 ± 1.40	4.1 × 10 ⁶
	NaM	nd ^b	nd ^b	<1.0 × 10 ³
	A	0.34 ± 0.17	23.8 ± 12.7	1.4 × 10 ⁴
	G	nd ^b	nd ^b	<1.0 × 10 ³
	C	6.29 ± 0.83	21.0 ± 6.1	3.0 × 10 ⁵
	T	5.66 ± 0.28	6.85 ± 0.80	8.3 × 10 ⁵

^a See Experimental Section for experimental details. ^b Rates too slow to independently determine k_{cat} and K_M .

Table S9. Kinetic data for insertion of **dNaM** opposite **d5SICS** in sequence context III^a

5'-d (TAATACGACTCACTATAGGGAGT)
3'-d (ATTATGCTGAGTGATATCCCTCAXTCTAGGTTACGGCAGGATCGC)

X(Template)	dYTP	k_{cat} (min ⁻¹)	K_M (μM)	k_{cat}/K_M (M ⁻¹ min ⁻¹)
5SICS	NaM	20.4 ± 4.41	3.44 ± 1.37	5.9 × 10 ⁶

^a See Experimental Section for experimental details.

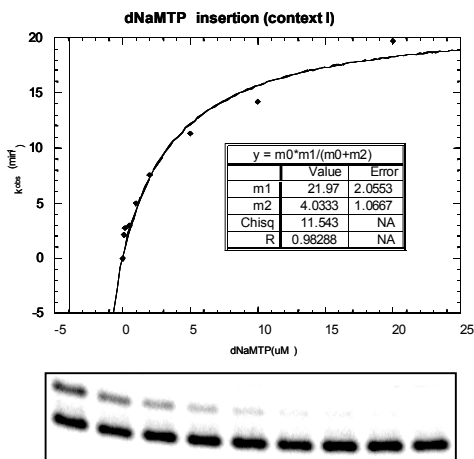
Table S10 Kinetic data for insertion of **dNaM** opposite **d5SICS** in sequence context IV^a

5'-d (TAATACGACTCACTATAGGGAGG)
3'-d (ATTATGCTGAGTGATATCCCTCCXTCTAGGTTACGGCAGGATCGC)

X(Template)	dYTP	k_{cat} (min ⁻¹)	K_M (μM)	k_{cat}/K_M (M ⁻¹ min ⁻¹)
5SICS	NaM	29.0 ± 7.7	0.45 ± 0.06	6.4 × 10 ⁷

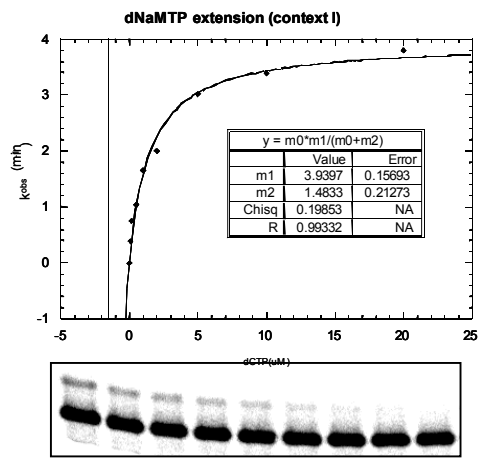
^a See Experimental Section for experimental details.

1) 5'-d(TAATACGACTCACTATAGGGAG Δ) ← dNaMTP
 3'-d(ATTATGCTGAGTGATATCCCTC $\underline{\text{I}}$ (d5SICS) $\underline{\text{G}}$ CTAGGTTACGGCAGGATCGC)



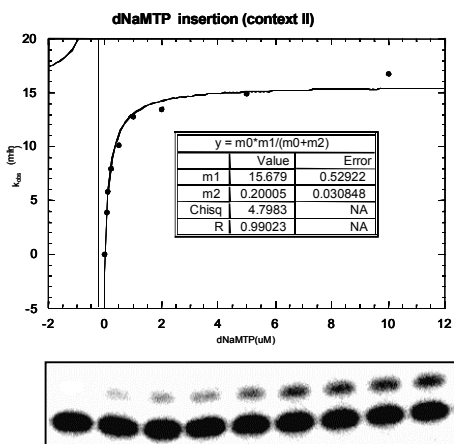
dNaMTP [μM] 20 10 5 2 1 0.5 0.2 0.1 0.05

2) 5'-d(TAATACGACTCACTATAGGGAGdNaM) ← dCTP
 3'-d(ATTATGCTGAGTGATATCCCTC $\underline{\text{I}}$ (d5SICS) $\underline{\text{G}}$ CTAGGTTACGGCAGGATCGC)



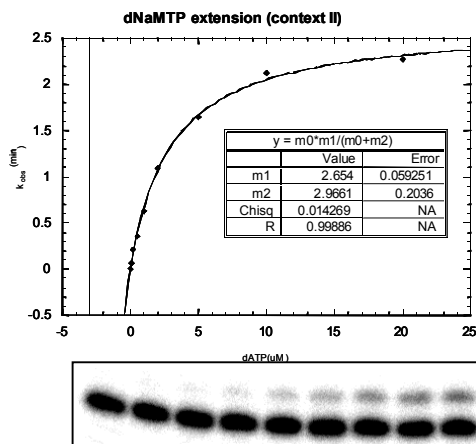
dCTP [μM] 20 10 5 2 1 0.5 0.2 0.1 0.05

3) 5'-d(TAATACGACTCACTATAGGGAG $\underline{\text{C}}$) ← dNaMTP
 3'-d(ATTATGCTGAGTGATATCCCTC $\underline{\text{G}}$ (d5SICS) $\underline{\text{I}}$ CTAGGTTACGGCAGGATCGC)



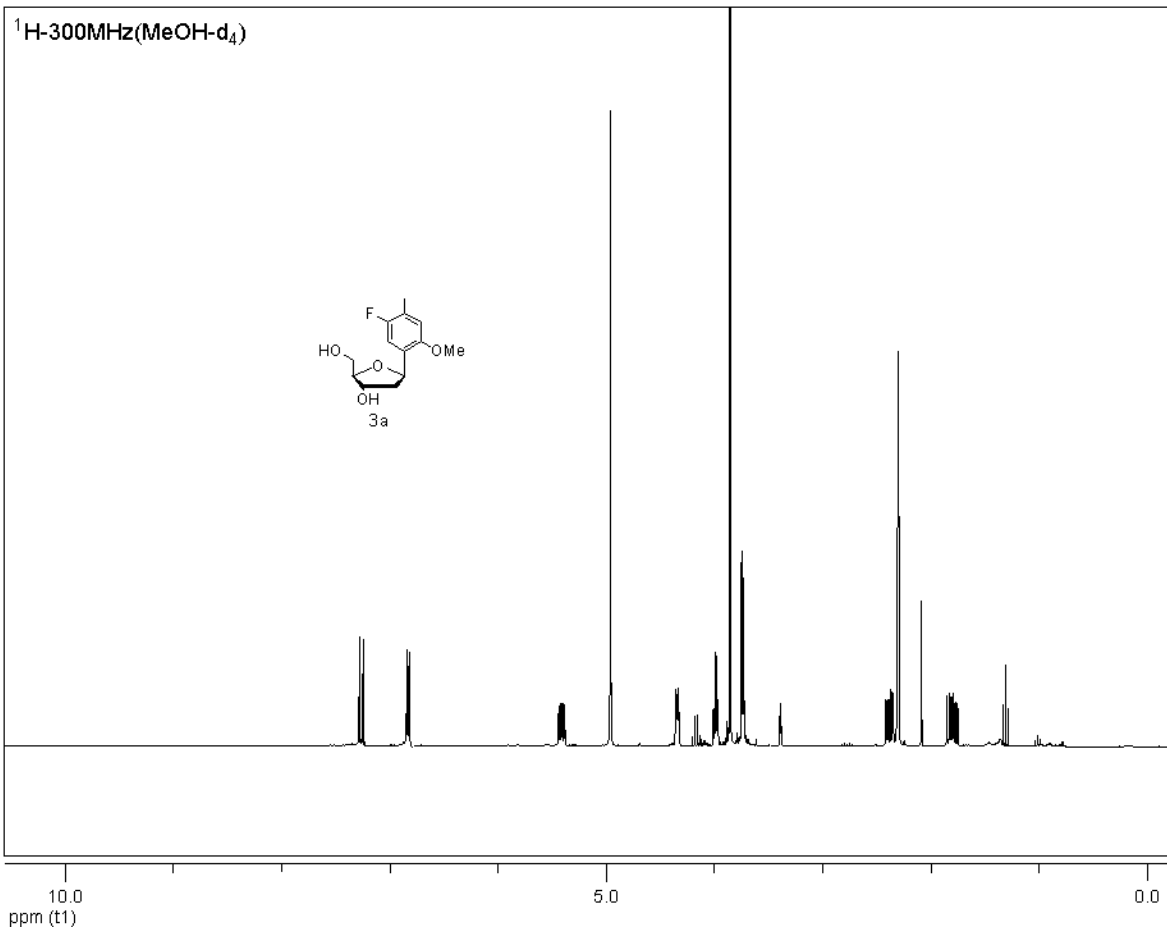
dNaMTP [μM] 0.02 0.05 0.1 0.2 0.5 1 2 5 10

4) 5'-d(TAATACGACTCACTATAGGGAGdNaM) ← dATP
 3'-d(ATTATGCTGAGTGATATCCCTC $\underline{\text{G}}$ (d5SICS) $\underline{\text{I}}$ CTAGGTTACGGCAGGATCGC)

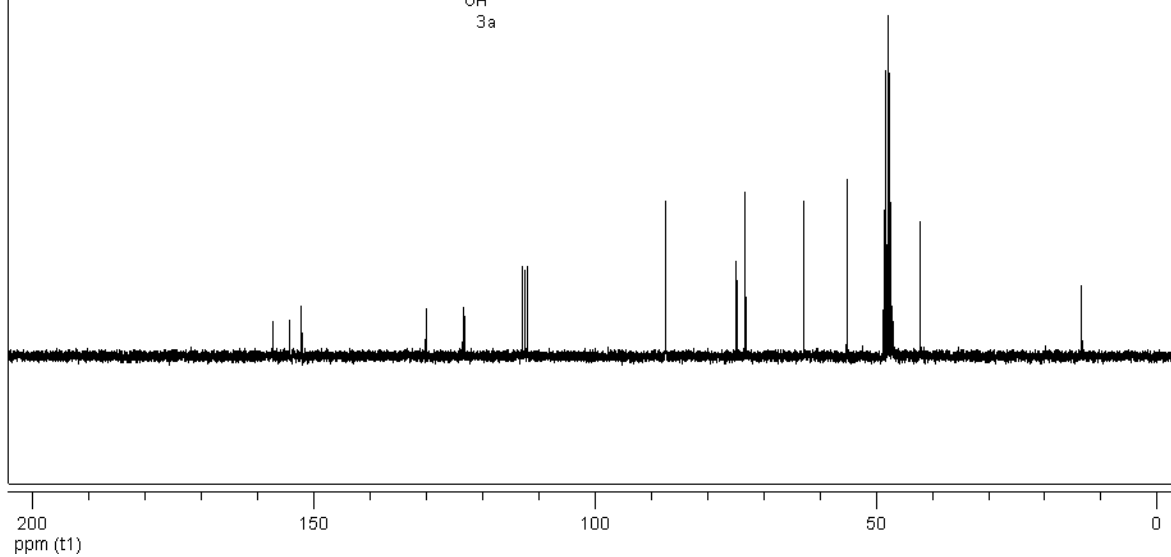
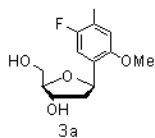


dATP [μM] 0.05 0.1 0.2 0.5 1 2 5 10 20

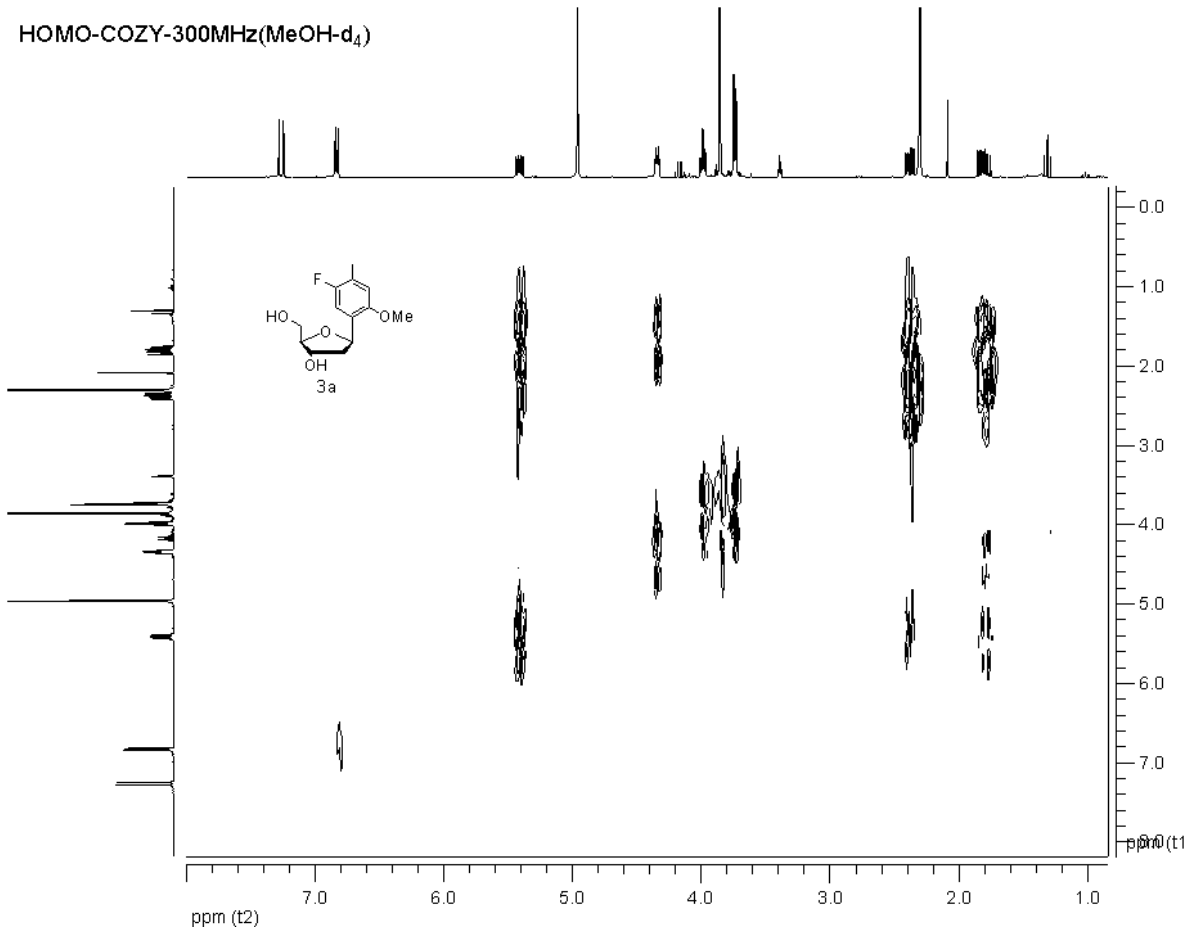
Figure S1. Raw gel data and plots of representative data (dNaMTP insertion opposite d5SICS and extension in sequence context I and II)

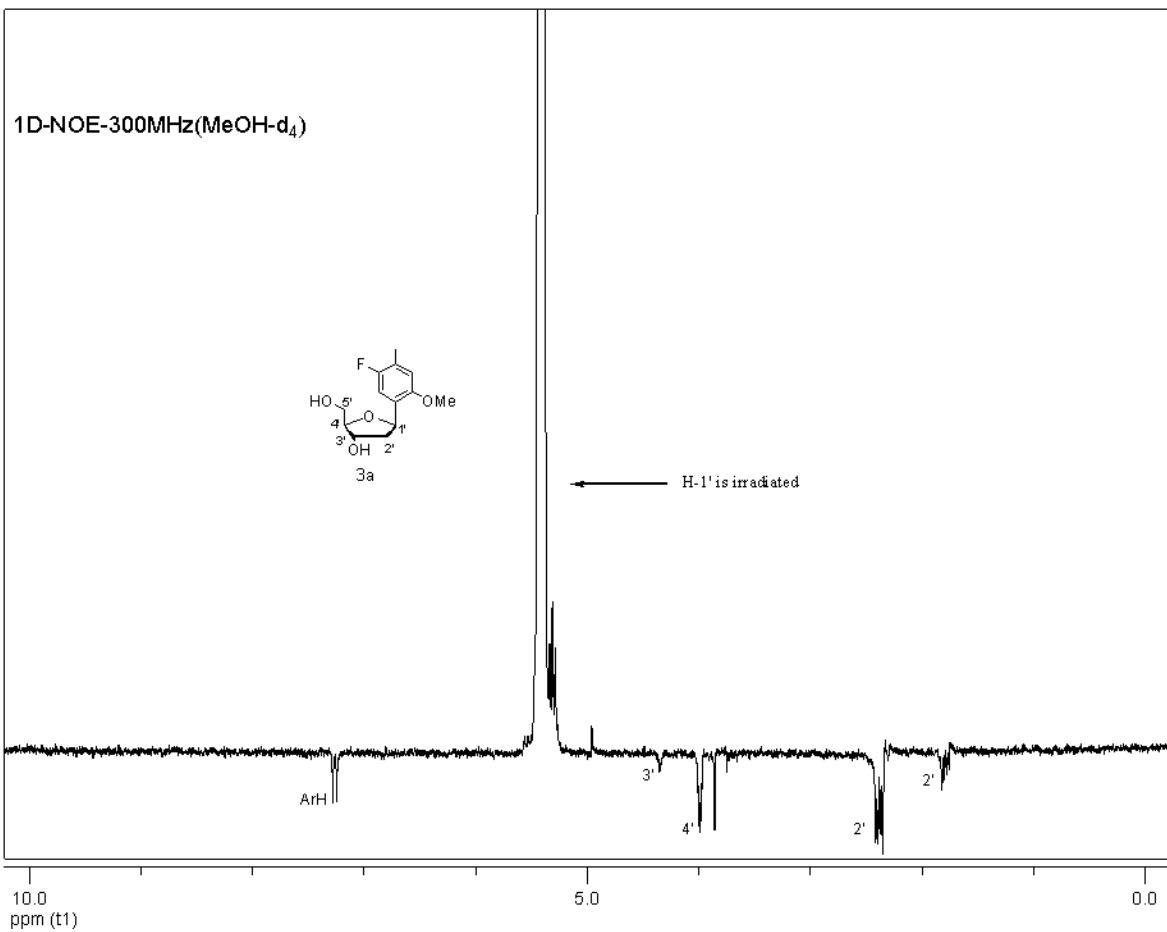


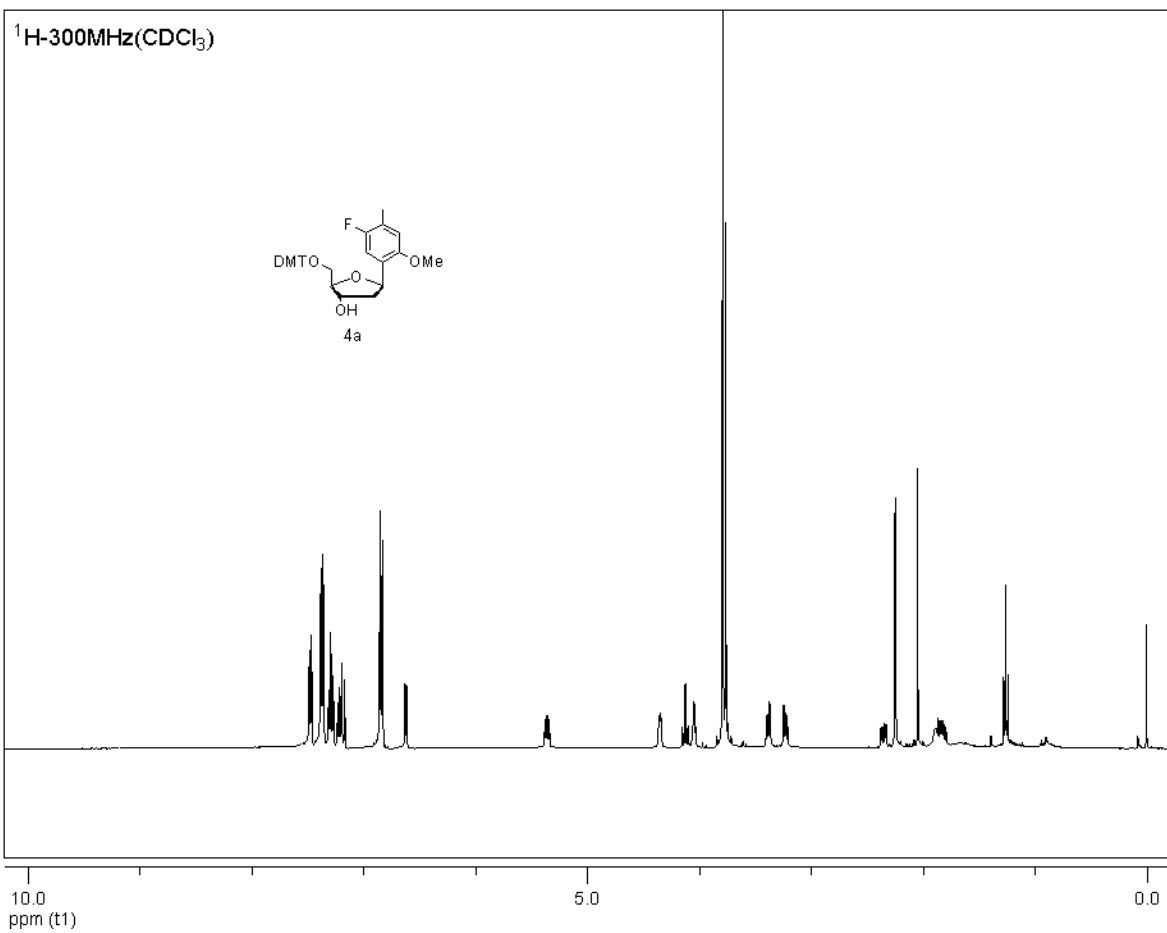
C¹³-300MHz(MeOH-d₄)



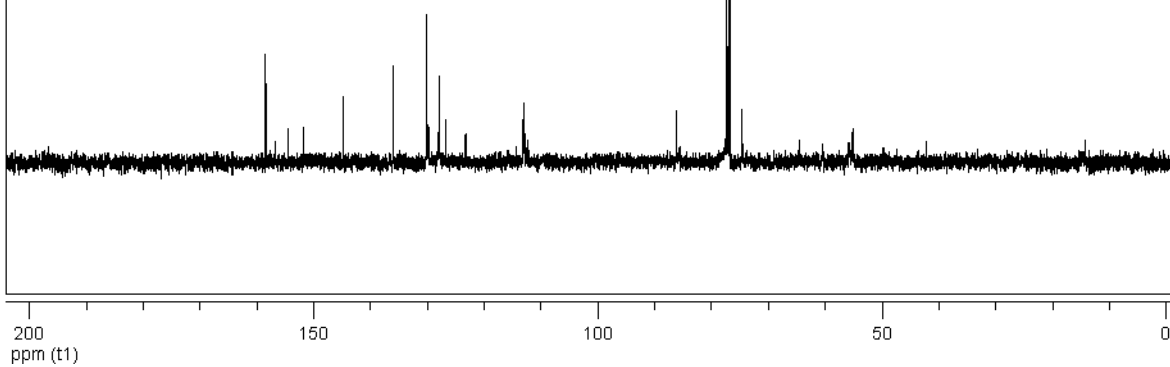
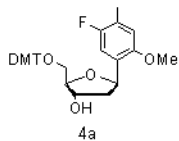
HOMO-COZY-300MHz(MeOH-d₄)

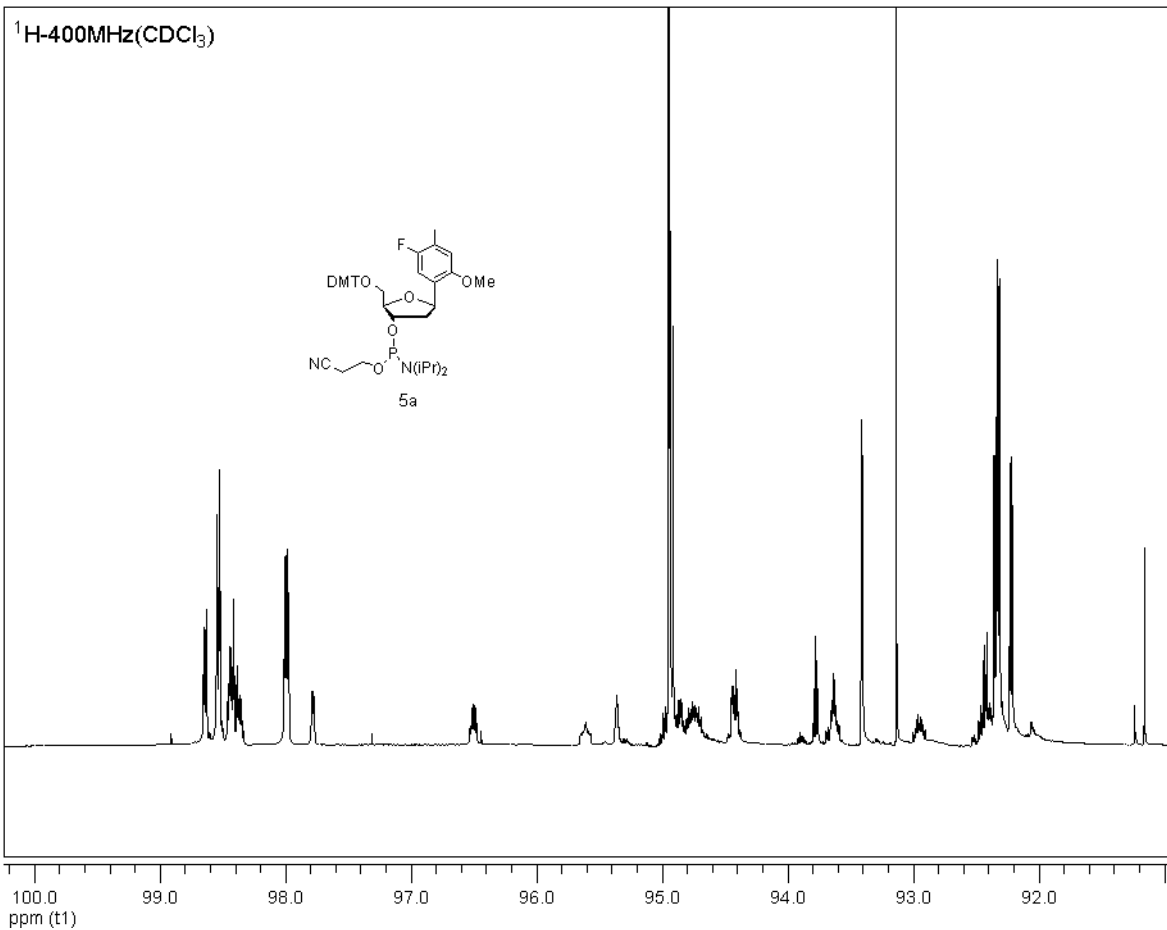




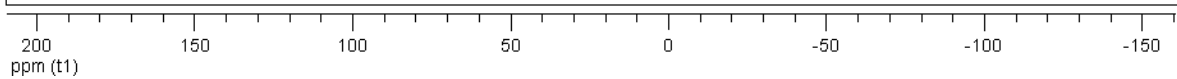
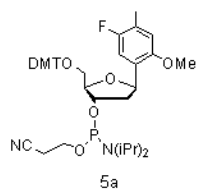


C¹³-300MHz(CDCl₃)

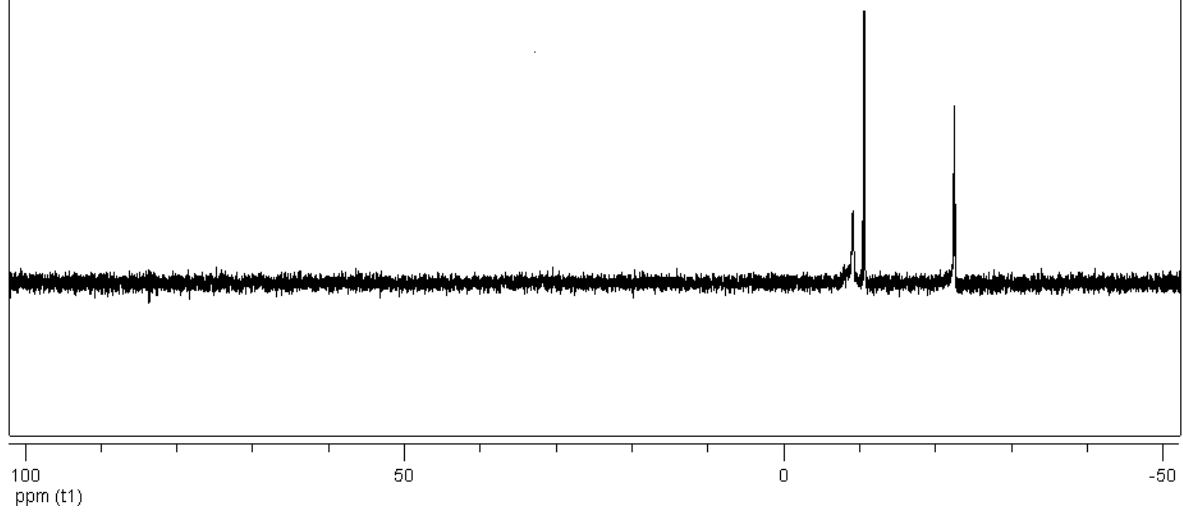
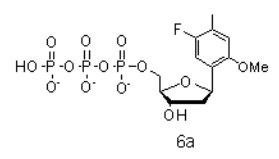


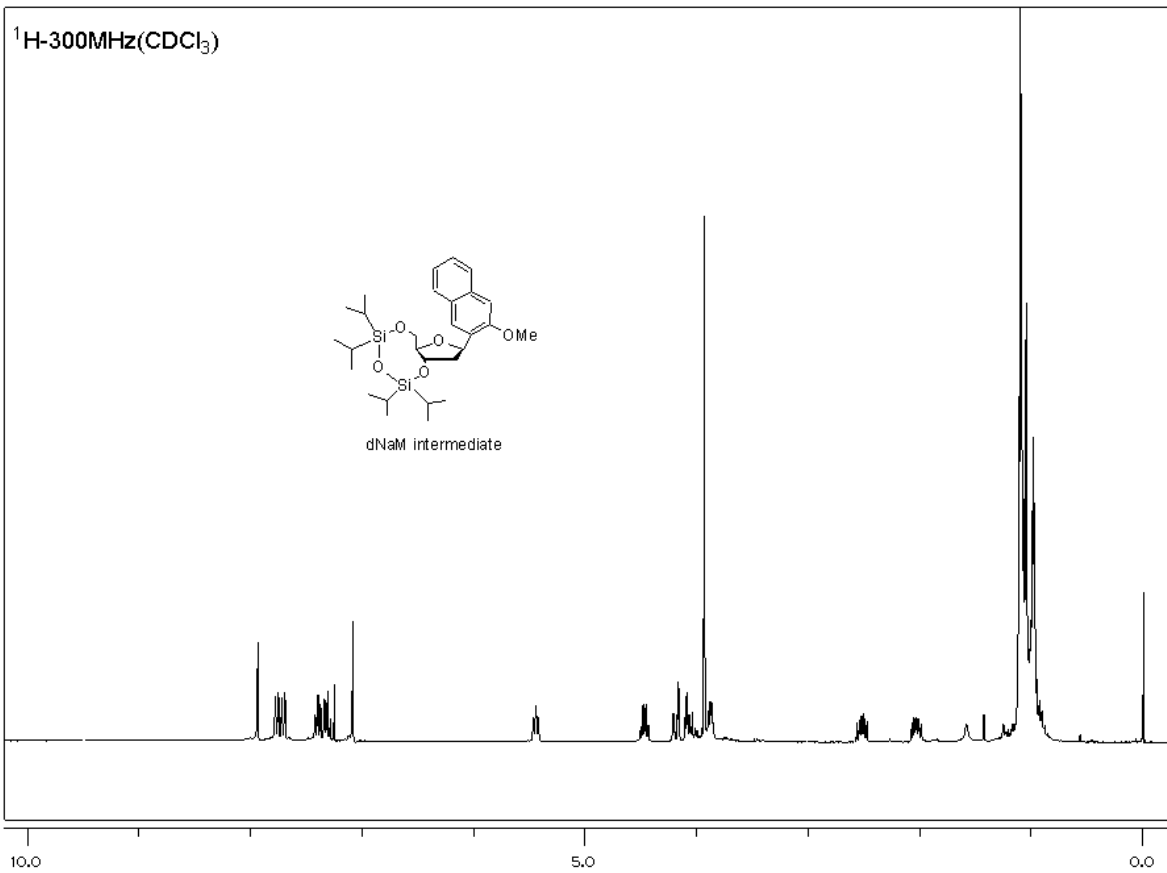


P³¹-400MHz(CDCl₃)

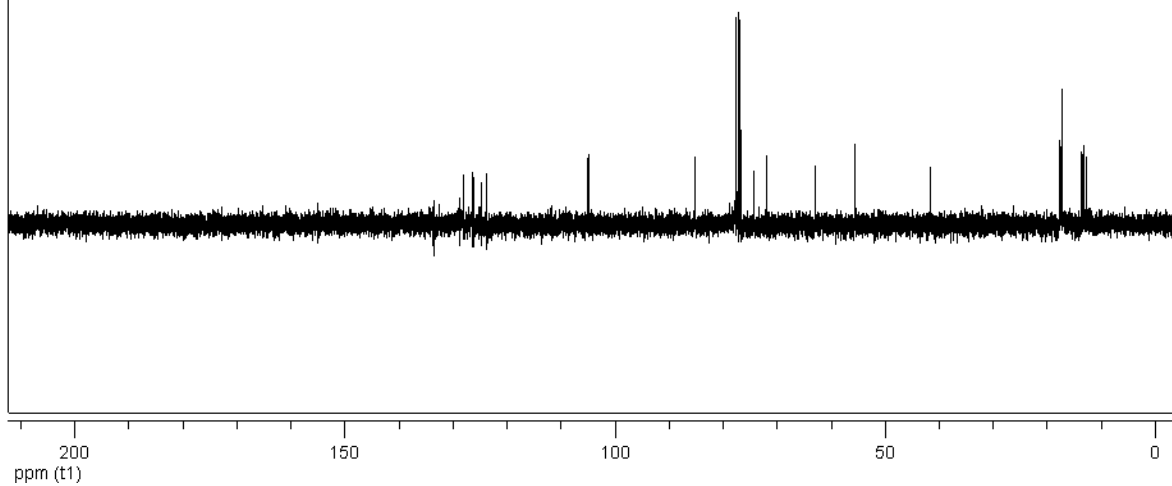
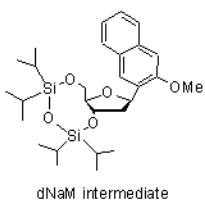


P³¹-400MHz(D₂O)

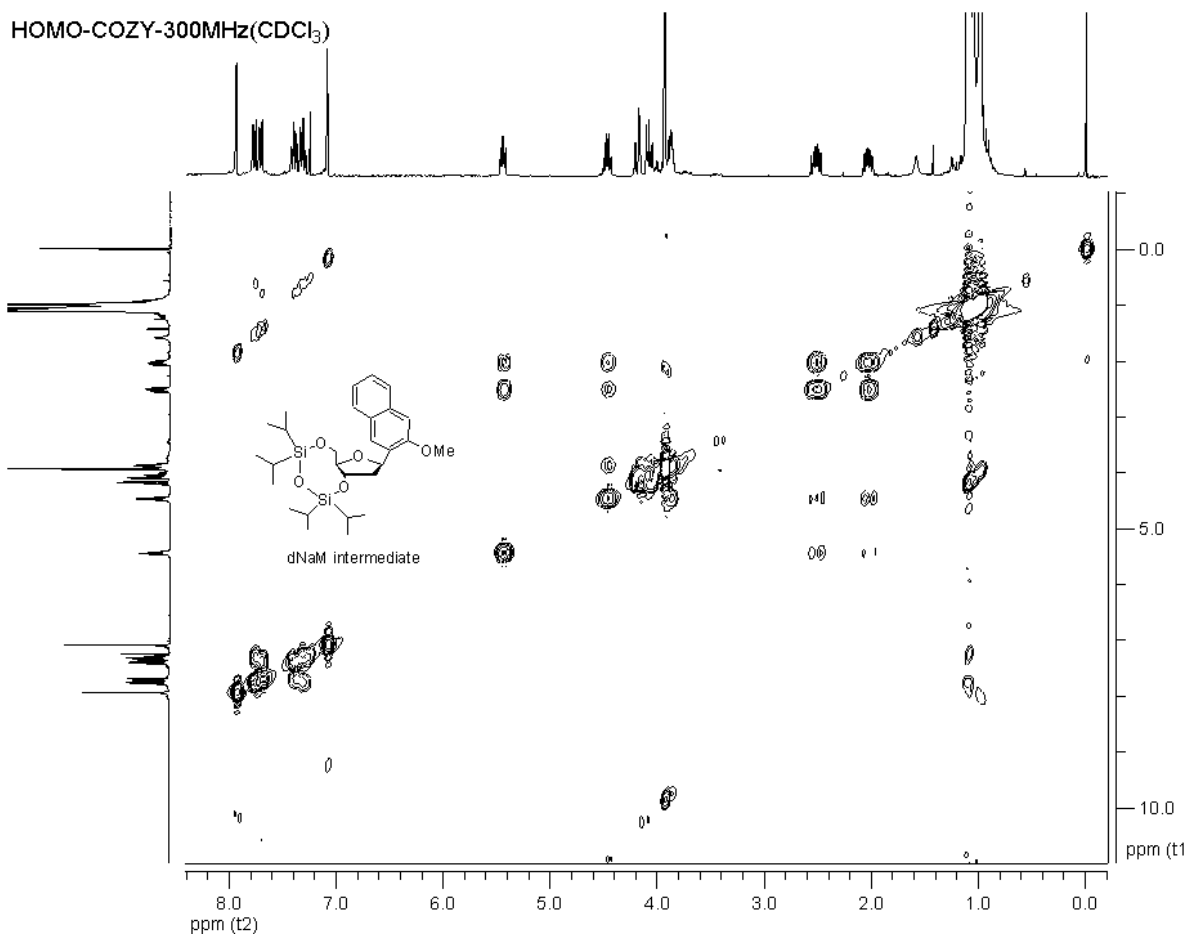


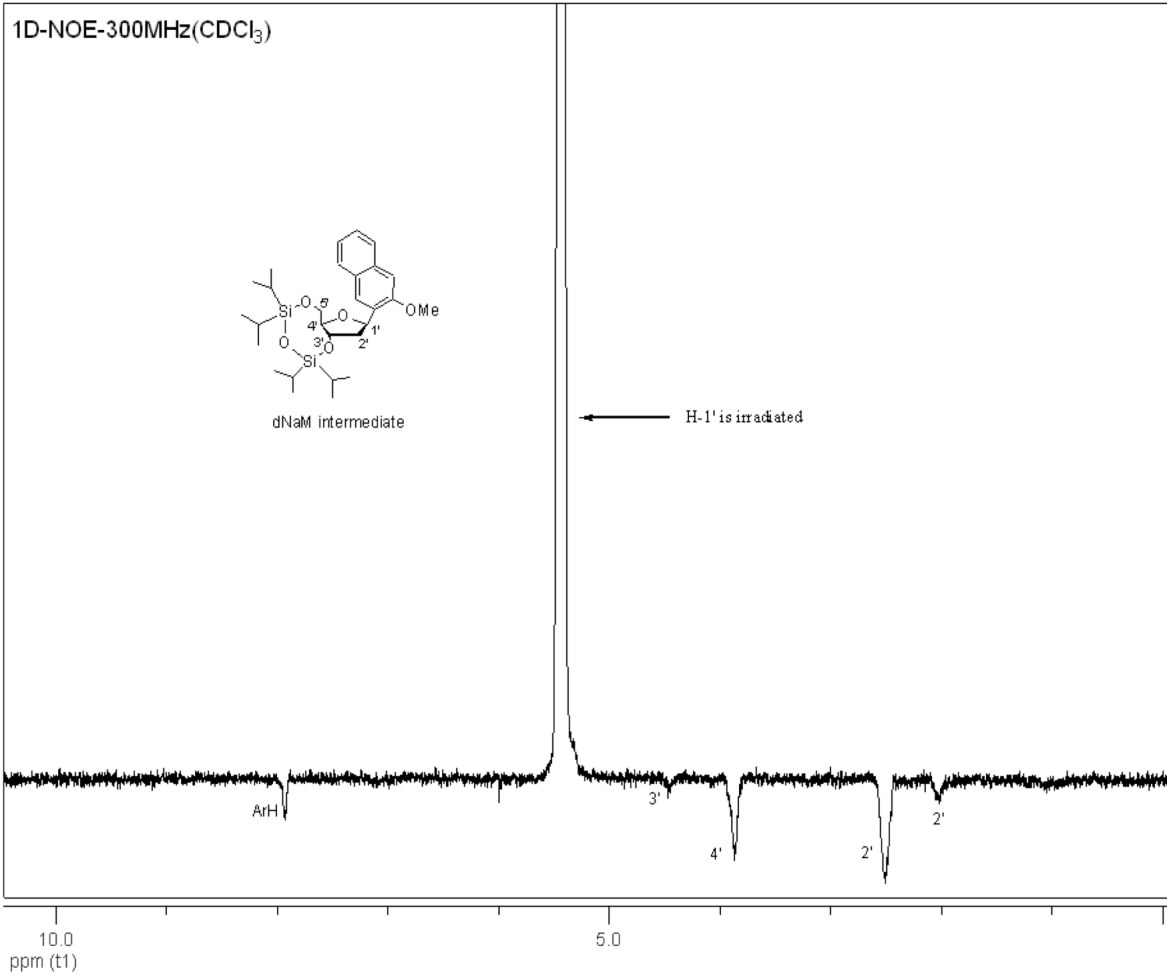


C¹³-300MHz(CDCl₃)

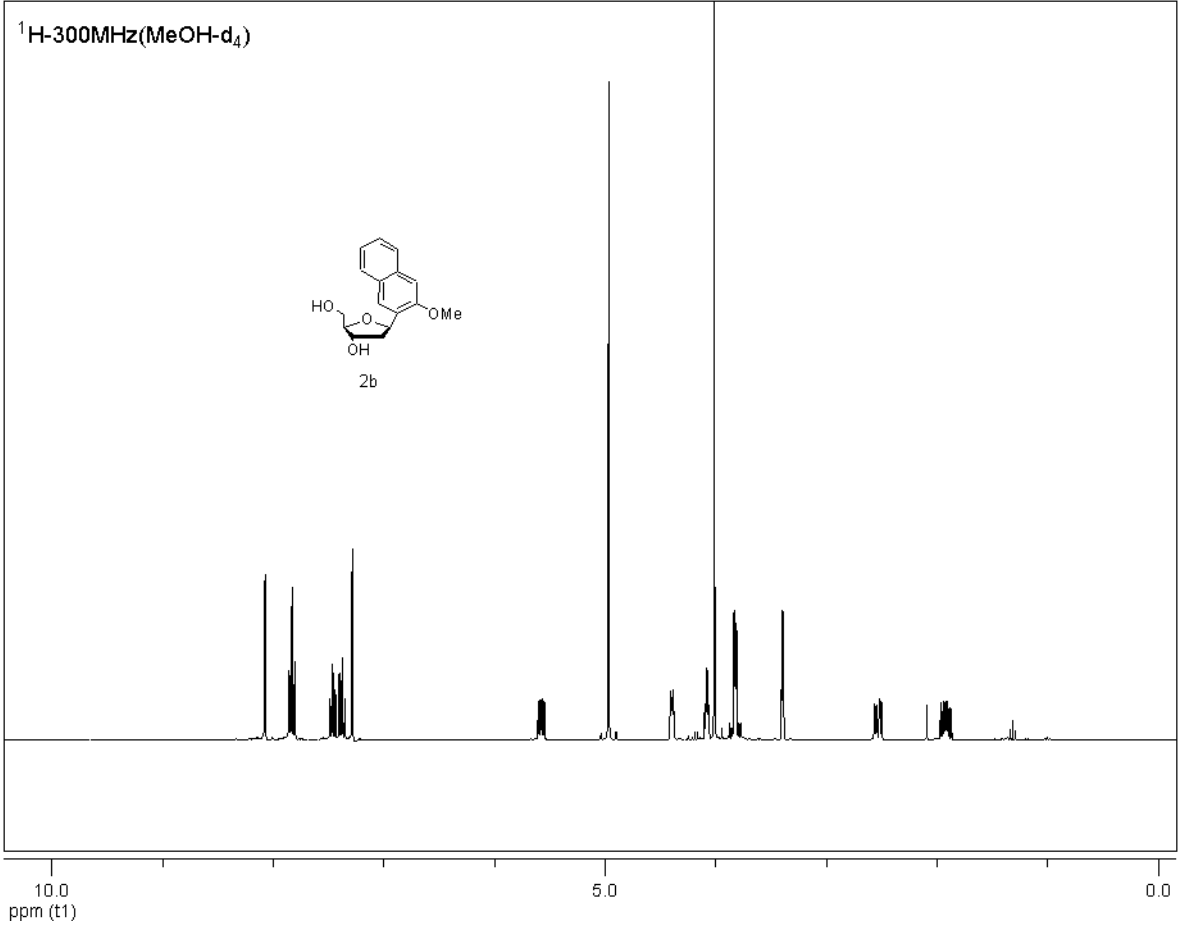
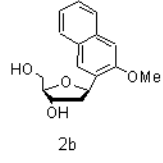


HOMO-COZY-300MHz(CDC₃)

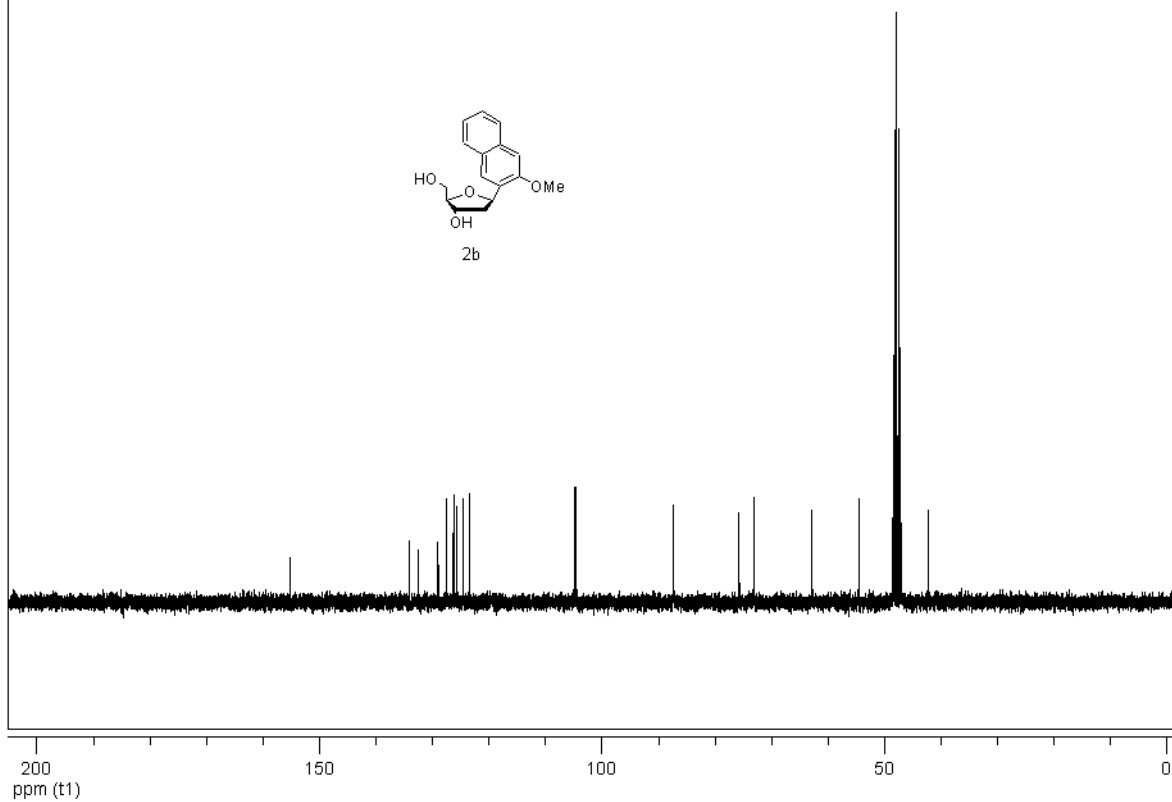
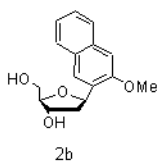


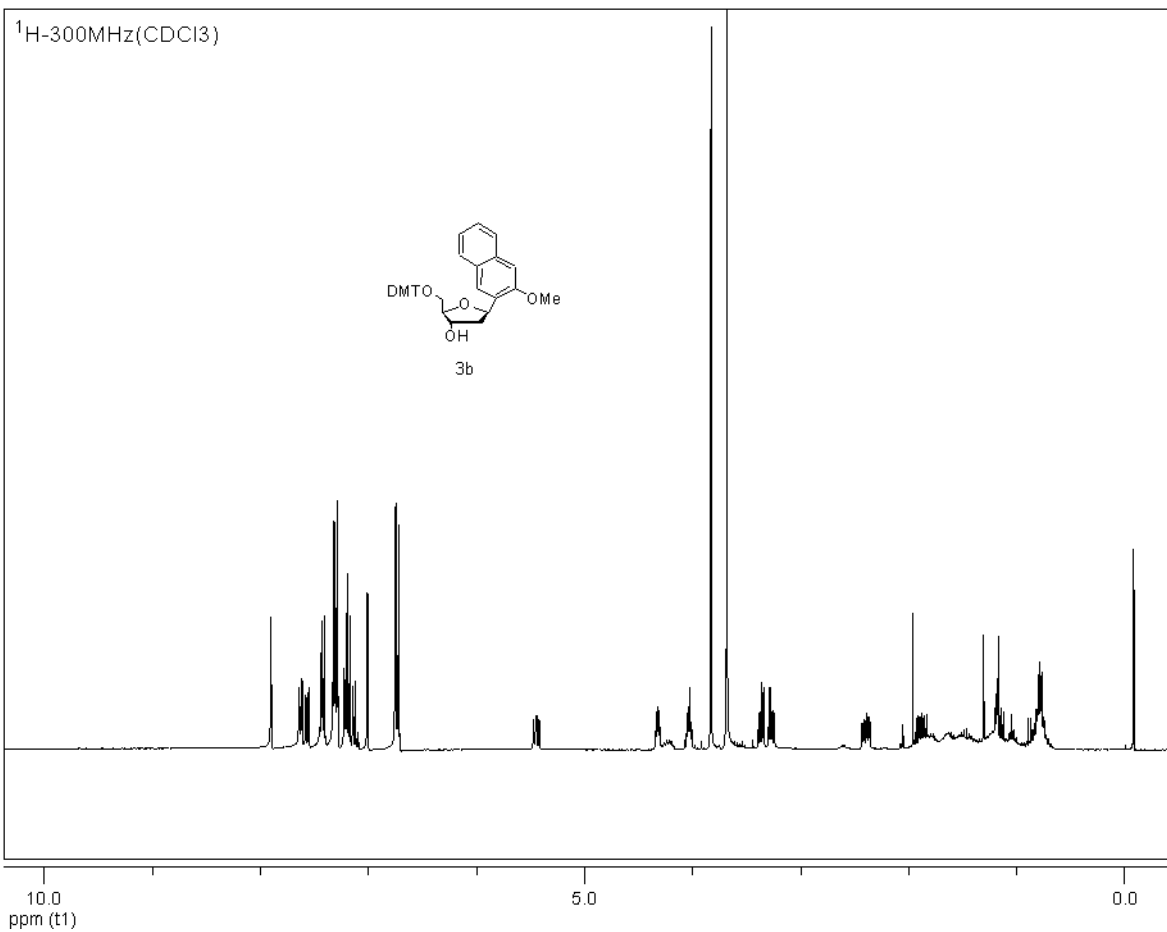


¹H-300MHz(MeOH-d₄)

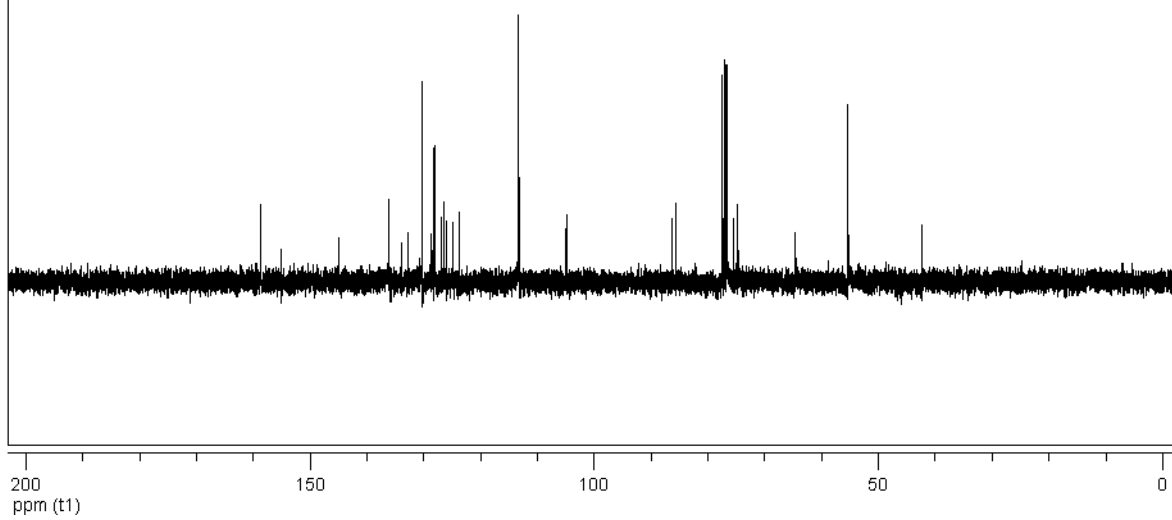
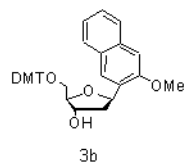


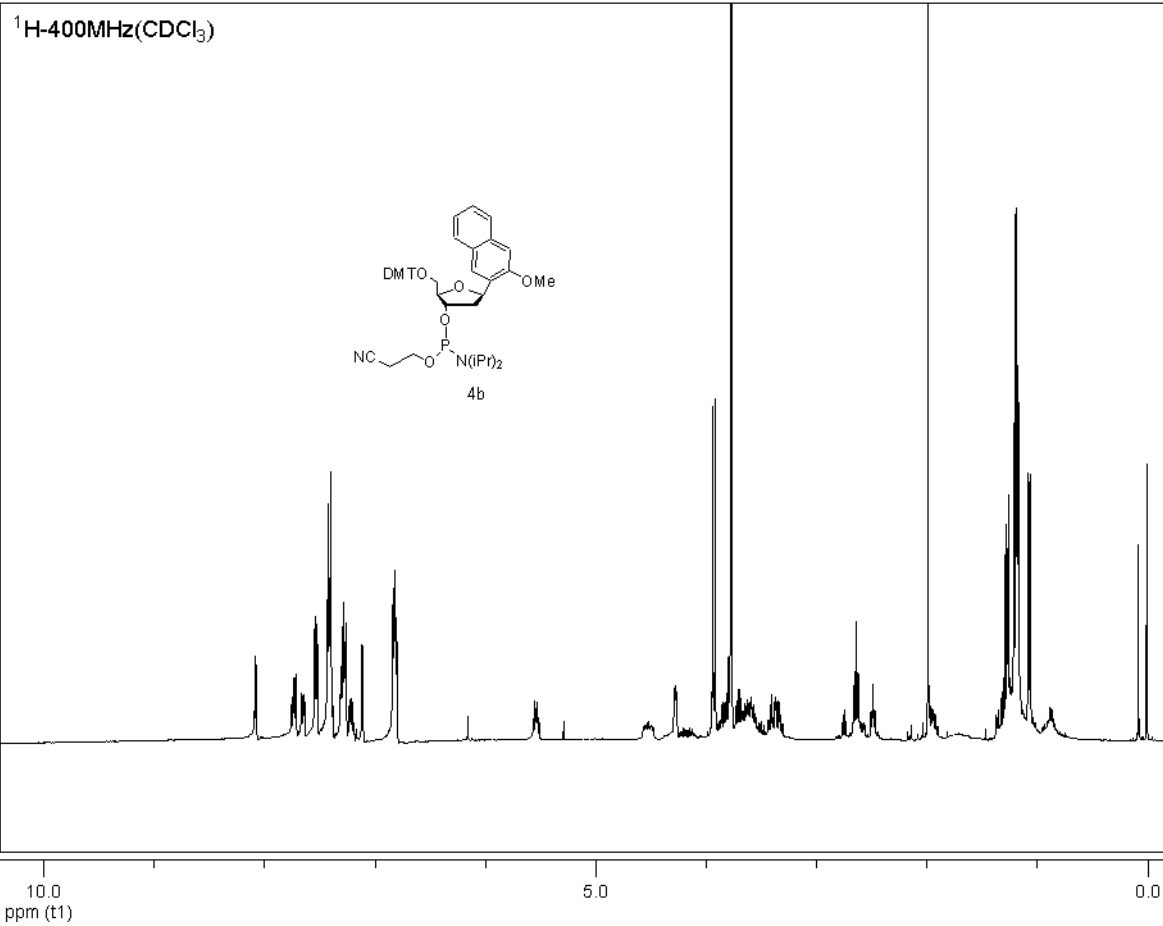
C¹³-300MHz(MeOH-d₄)



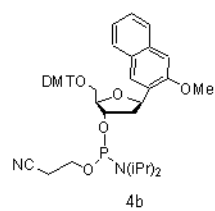


C¹³-300MHz(CDCl₃)





P³¹-400MHz(CDCl₃)



200
ppm (t1)

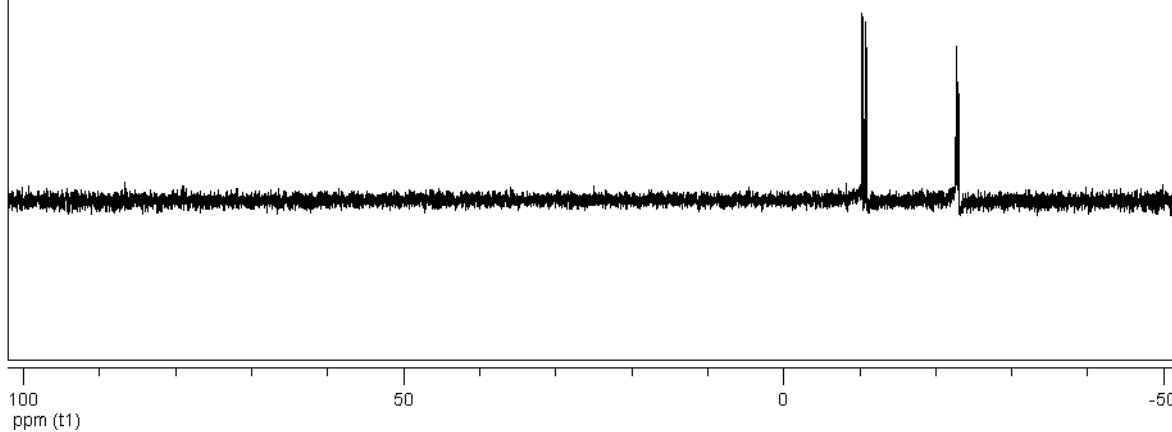
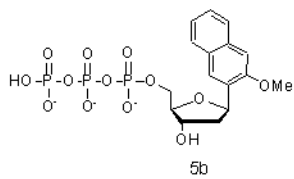
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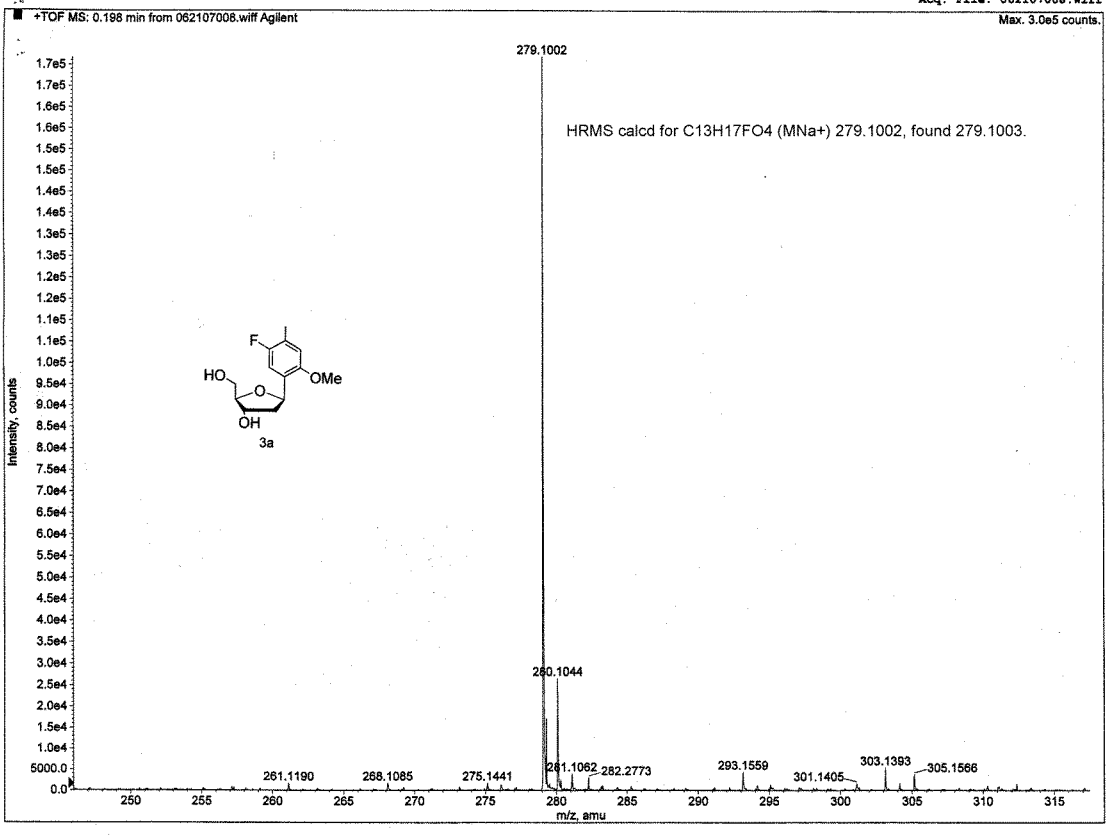
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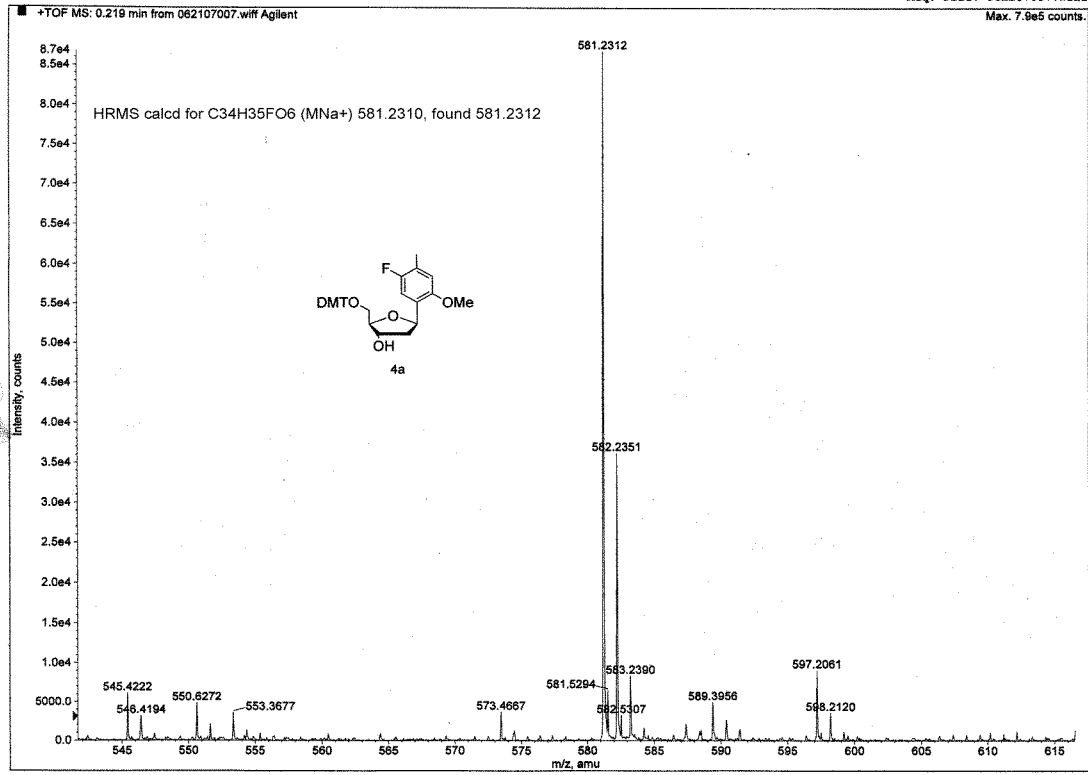
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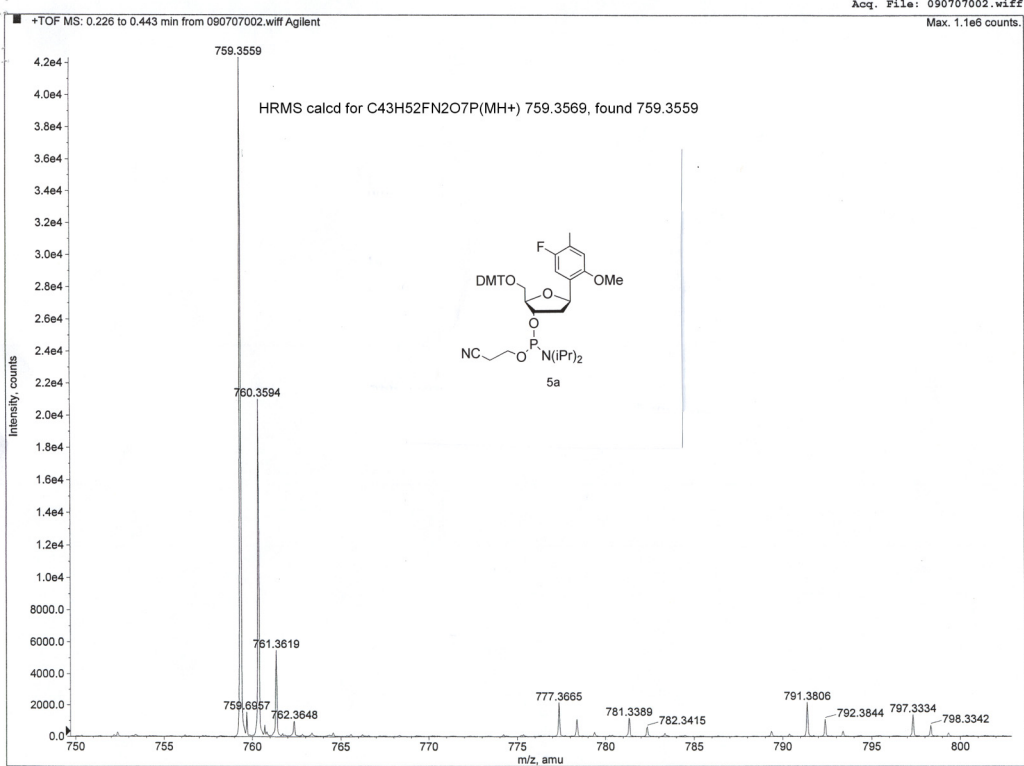
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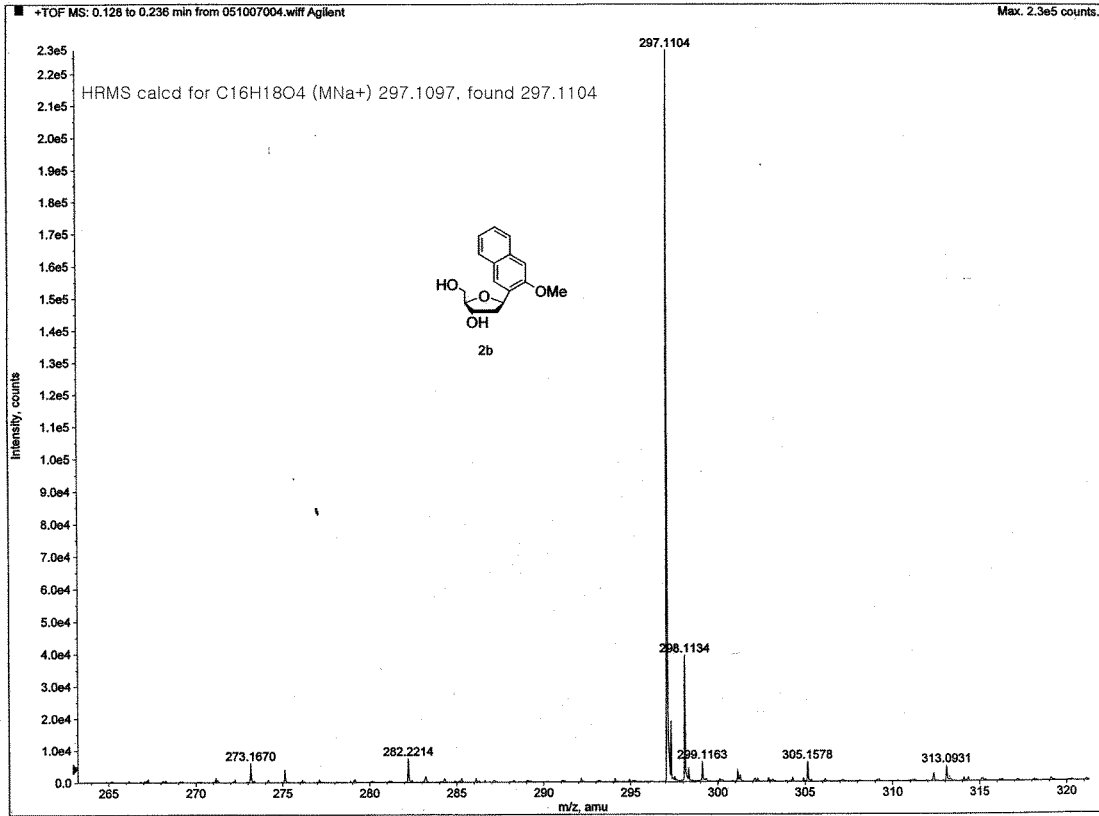
P³¹-400MHz(D₂O)

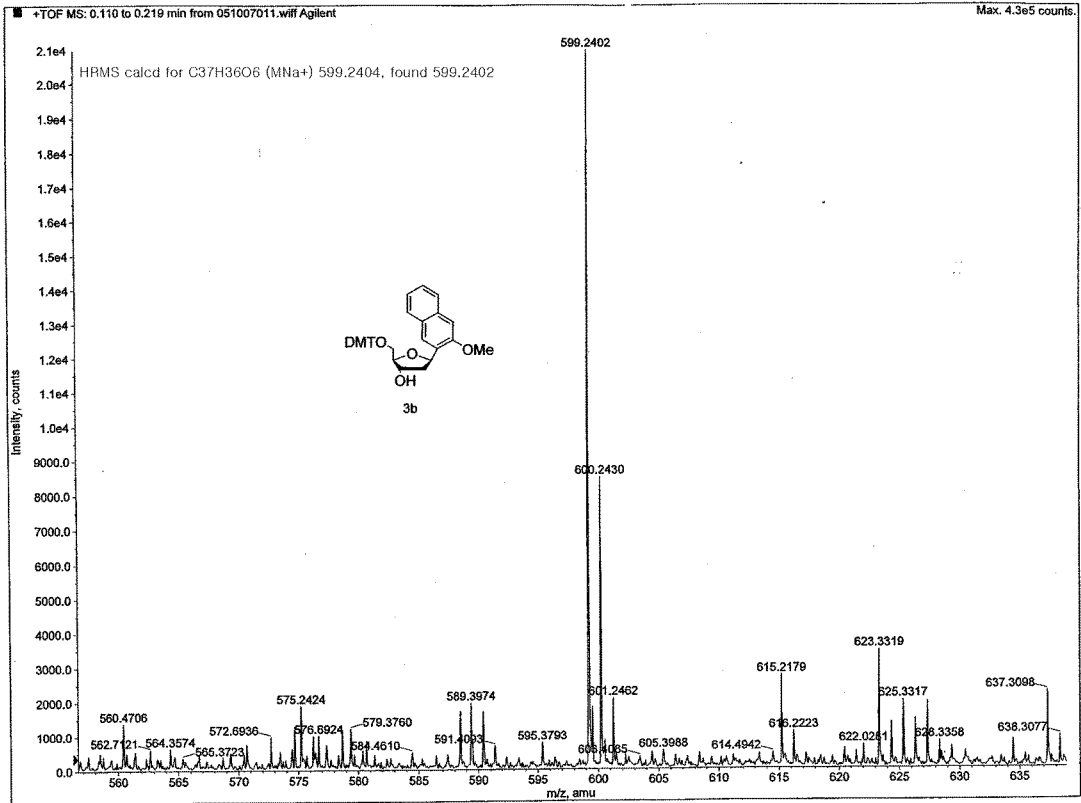












Polarity/Scan Type: Positive

Sample Name: xbya5757

Acq. I

Day, June 21, 2007

File: 062107004.wiff

Max. 4.3e5 counts.

