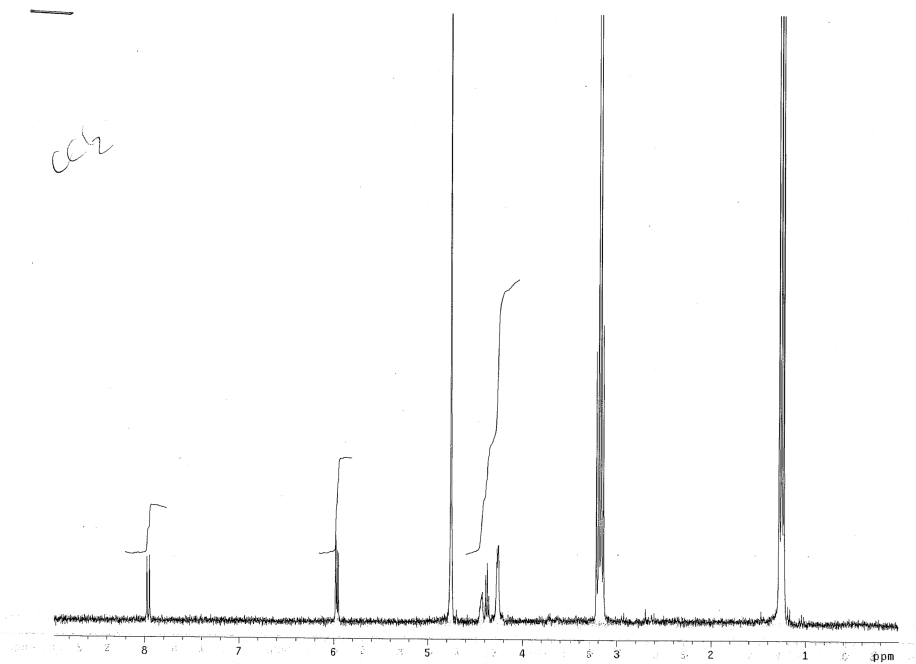
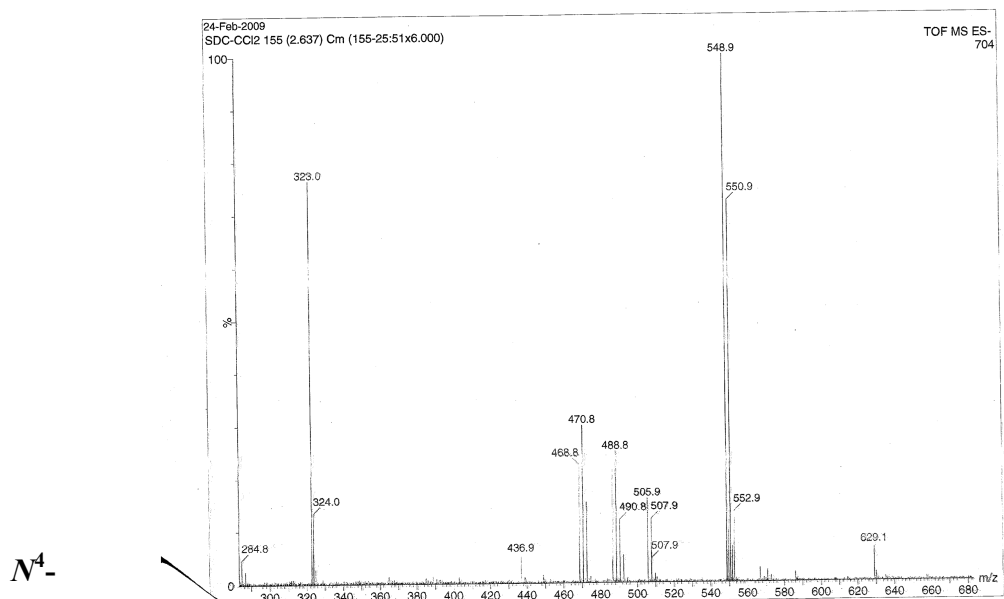


Supporting information

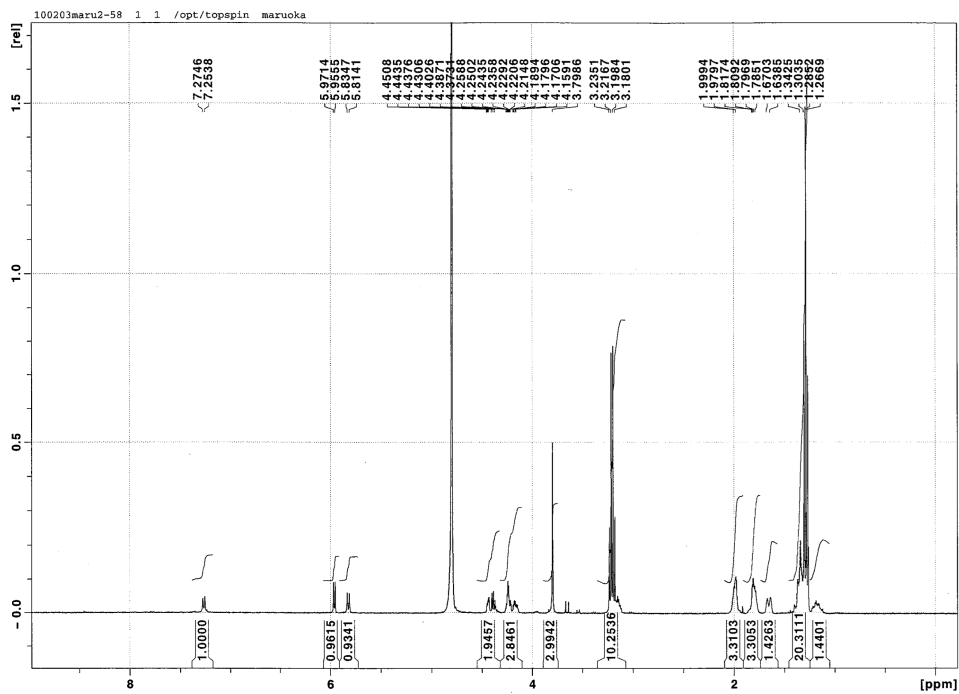
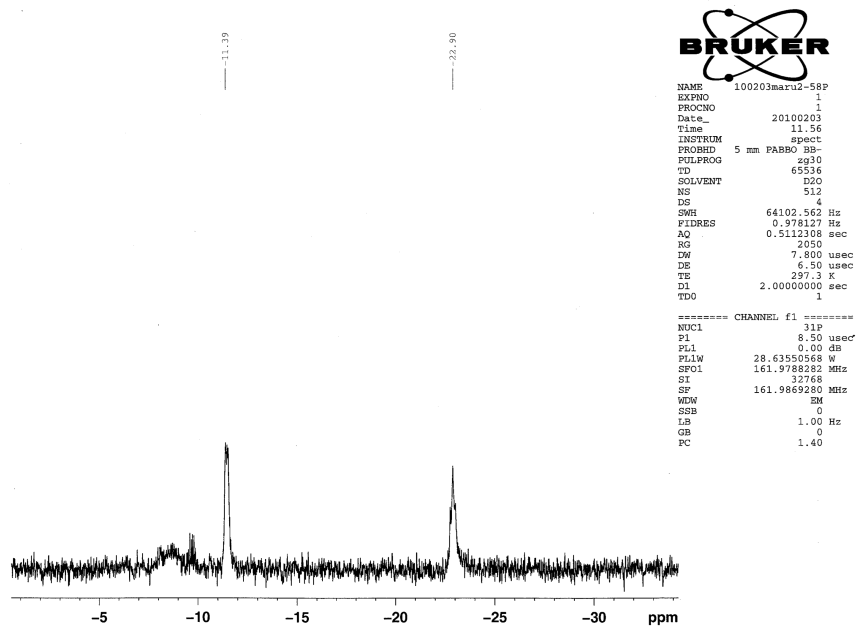
Pyrimidine Nucleotides with 4-Alkyloxyimino and Terminal Phosphate Ester Modifications as Selective Agonists of the P2Y₄ Receptor

Hiroshi Maruoka, M.P. Suresh Jayasekara, Matthew O. Barrett, Derek A. Frankin, Sonia de Castro, Nathaniel Kim, Stefano Costanzi, T. Kendall Harden, and Kenneth A. Jacobson

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Table 1. Crystal data and structure refinement for <i>N</i> ⁴ - <i>t</i> -BuO-cytosine.	S27
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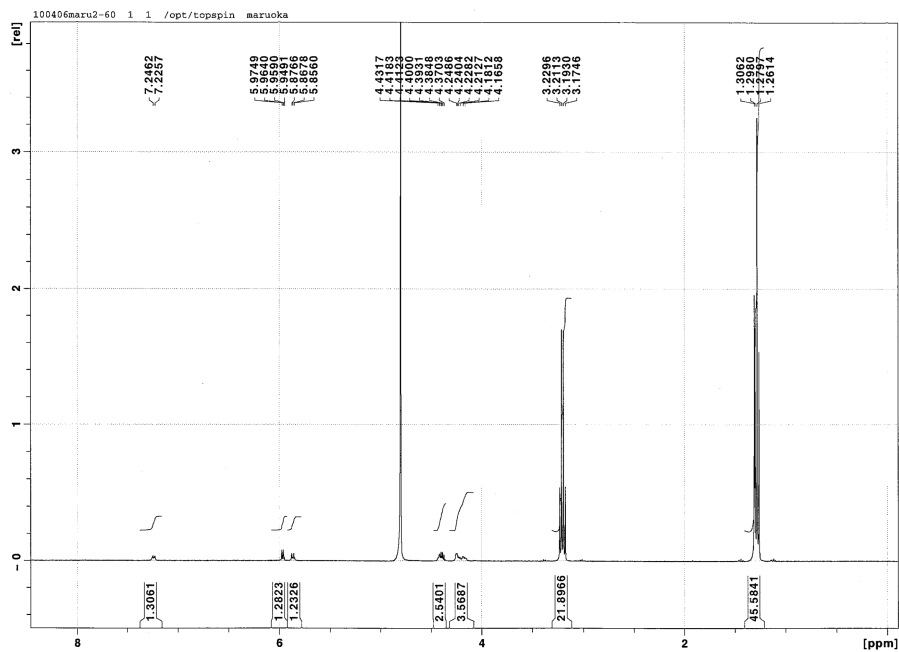
Uridine-5'- β,γ -dichloromethylene triphosphate ammonium salt (10) **^1H NMR****Mass**

Methoxycytidine-5'-triphosphate triethylammonium salt (11)

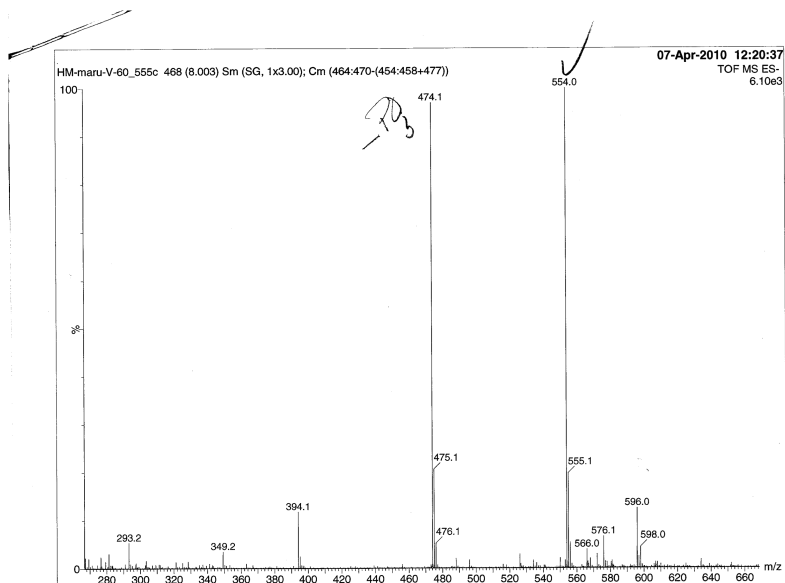
 ^1H NMR ^{31}P NMR

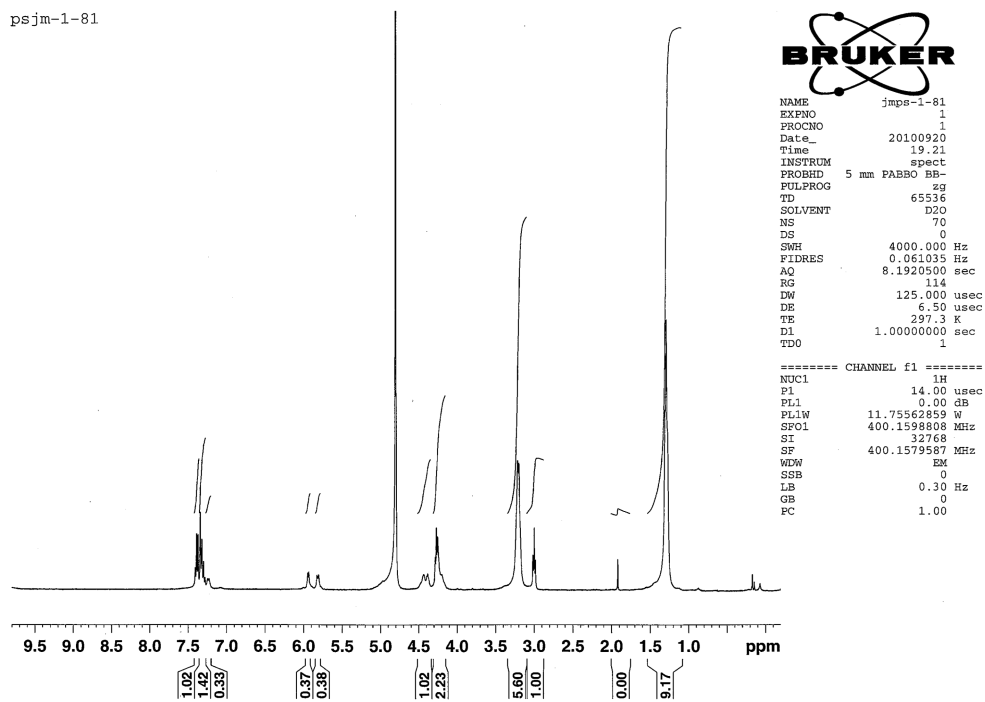
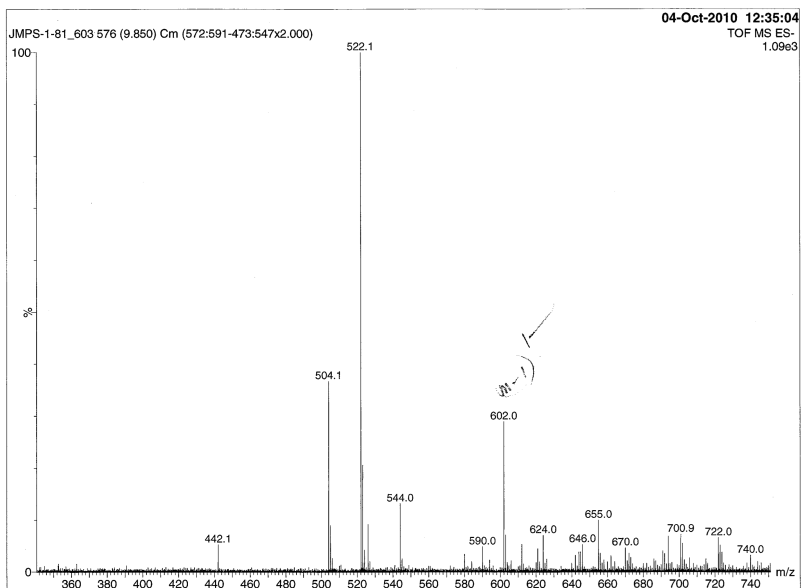
*N*⁴-*tert*-Butyloxycytidine-5'-triphosphate triethylammonium salt (13)

¹H NMR



Mass

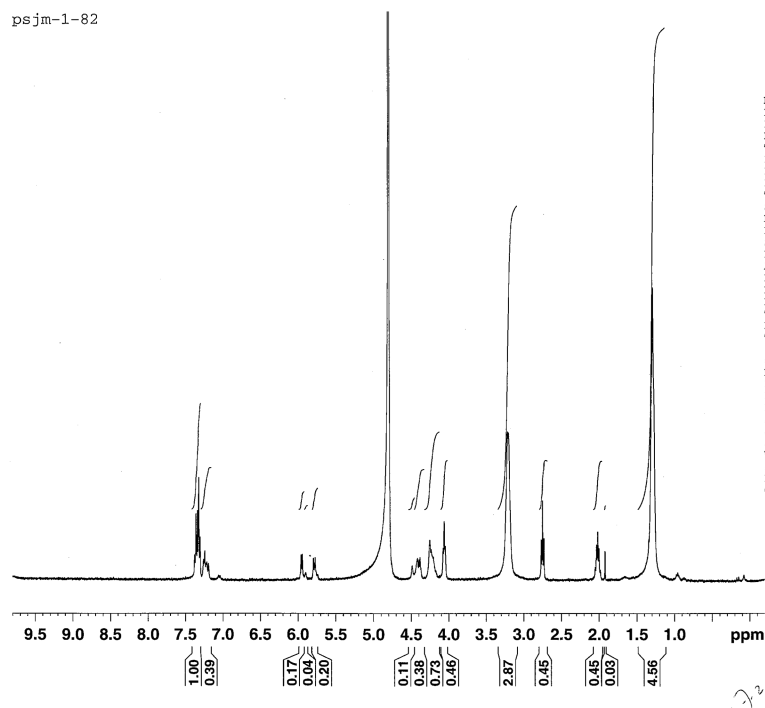


N*⁴-Phenylethoxycytidine-5'-triphosphate triethylammonium salt (15)*¹H NMR****Mass**

N^4 -Phenylpropoxycytidine-5'-triphosphate triethylammonium salt (16)

^1H NMR

psjm-1-82



```

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PROCNO        1
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PULPROG       zg
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SOLVENT       D2O
NS            27
DS            0
SWH           4000.000 Hz
FIDRES        0.061035 Hz
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DE            6.50 usec
TE            297.6 K
DL            1.0000000 sec
TDO           1

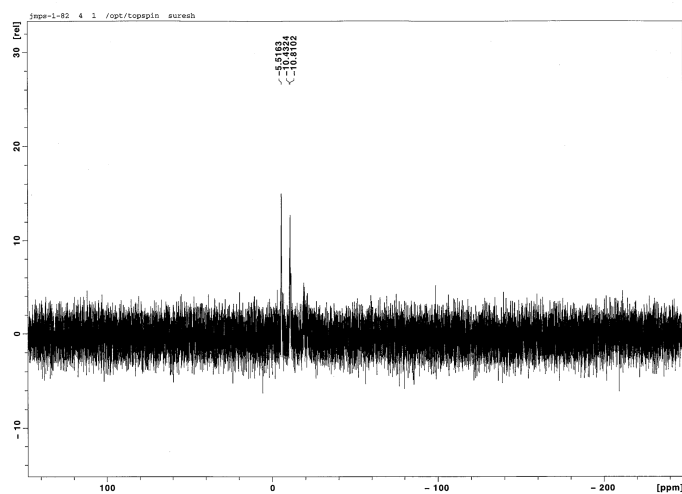
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GB            0
PC            1.00

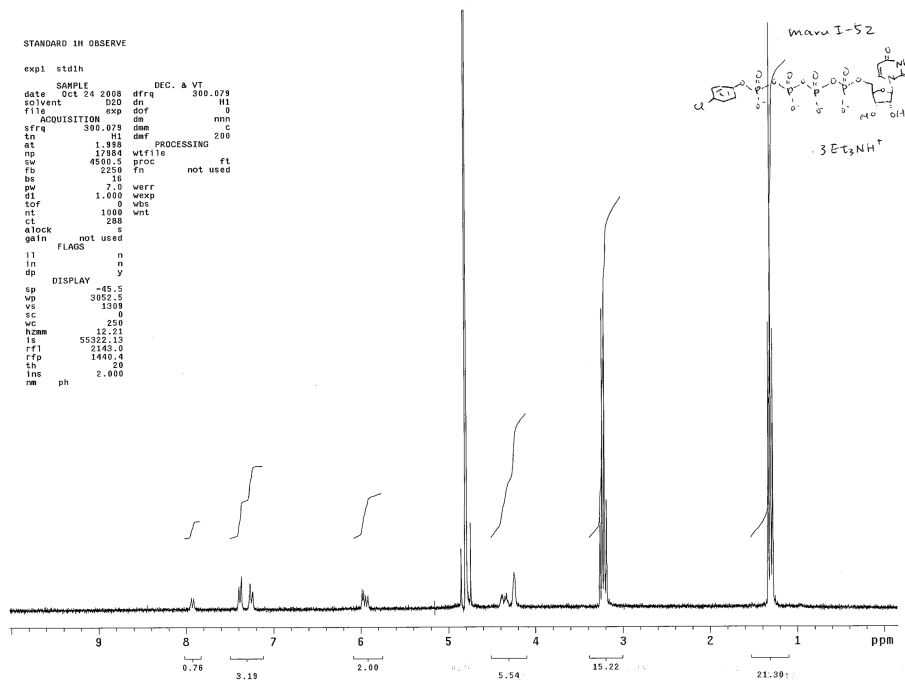
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^{31}P NMR

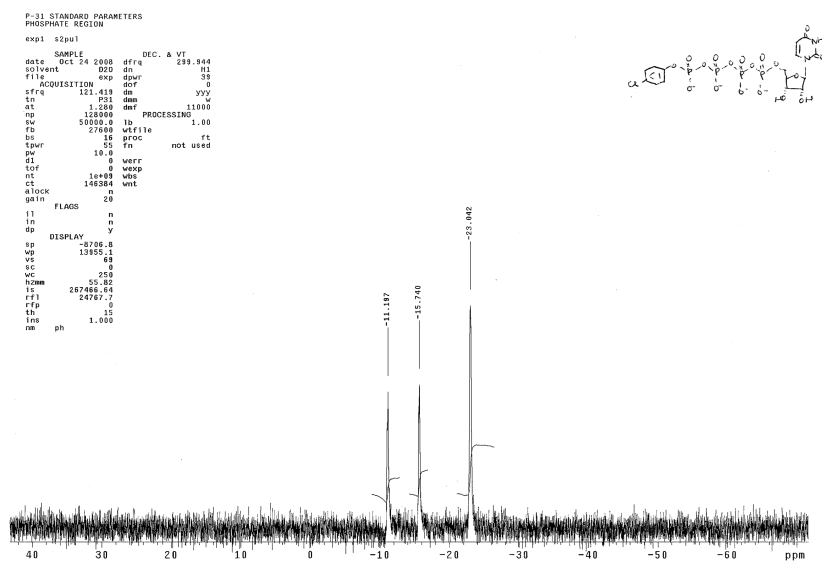


Uridine-5'-(4-chlorophenyl)- tetraphosphate triethylammonium salt (20)

^1H NMR

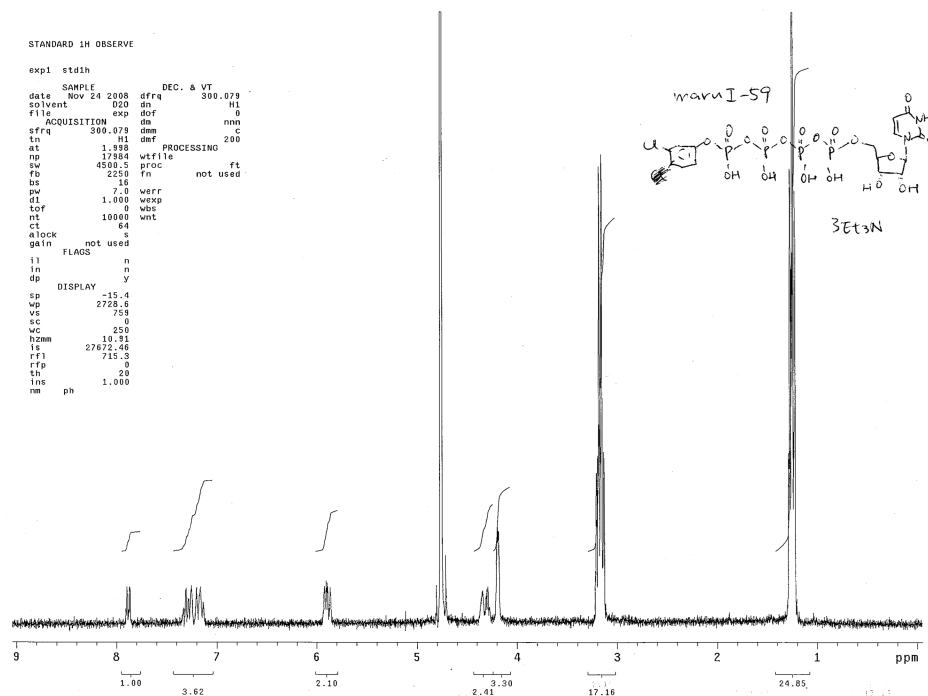


^{31}P NMR

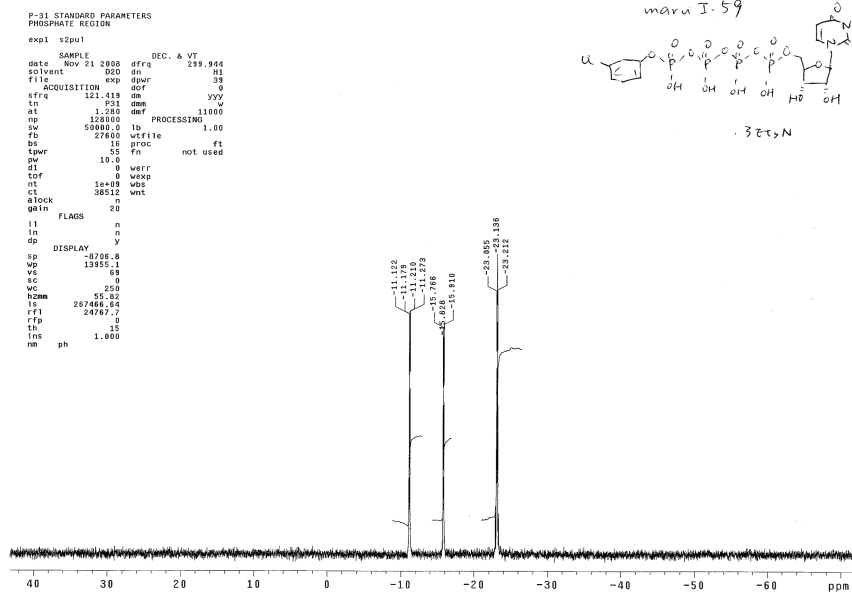


Uridine-5'-(3-chlorophenyl)- tetraphosphate triethylammonium salt (21)

^1H NMR

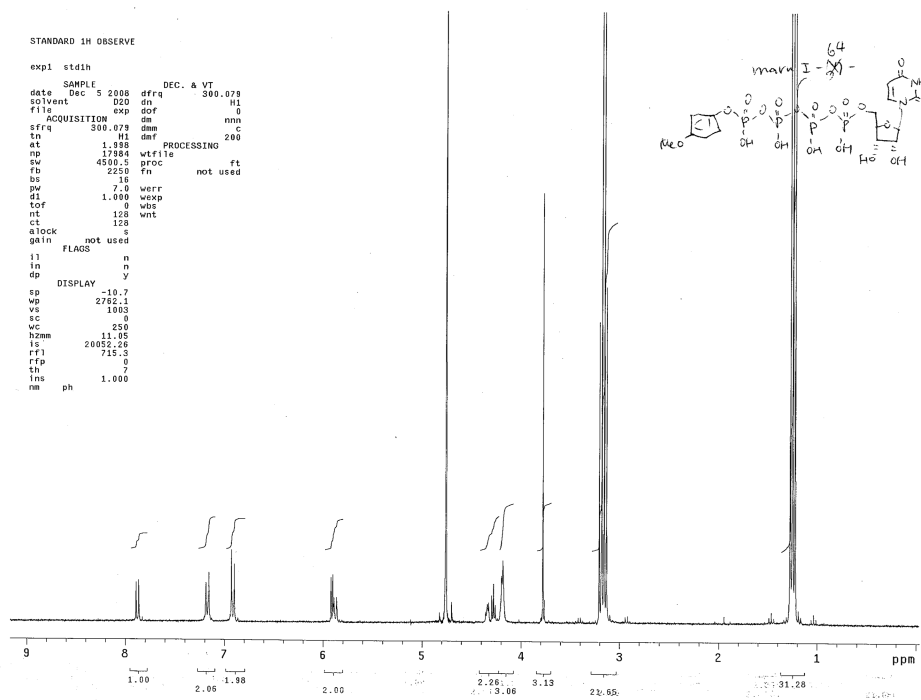


^{31}P NMR

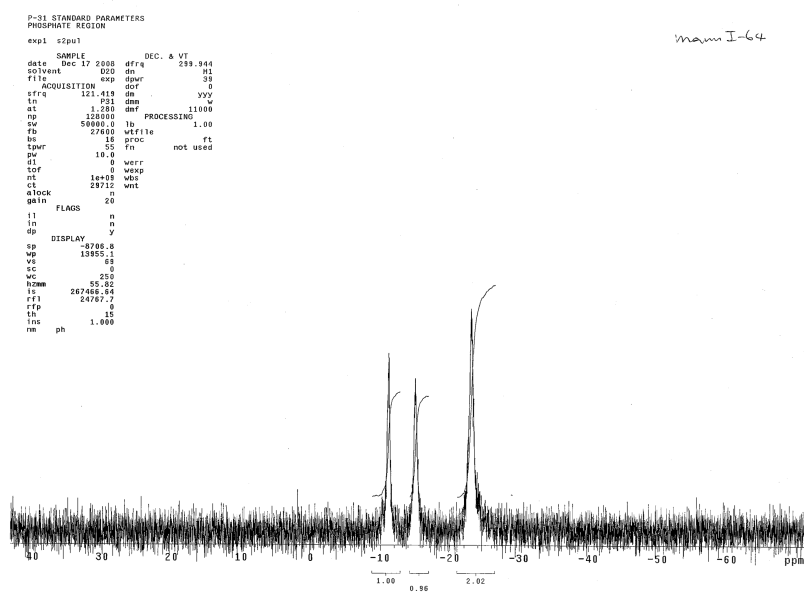


Uridine-5'-(4-methoxyphenyl)- tetraphosphate triethylammonium salt (22)

¹H NMR

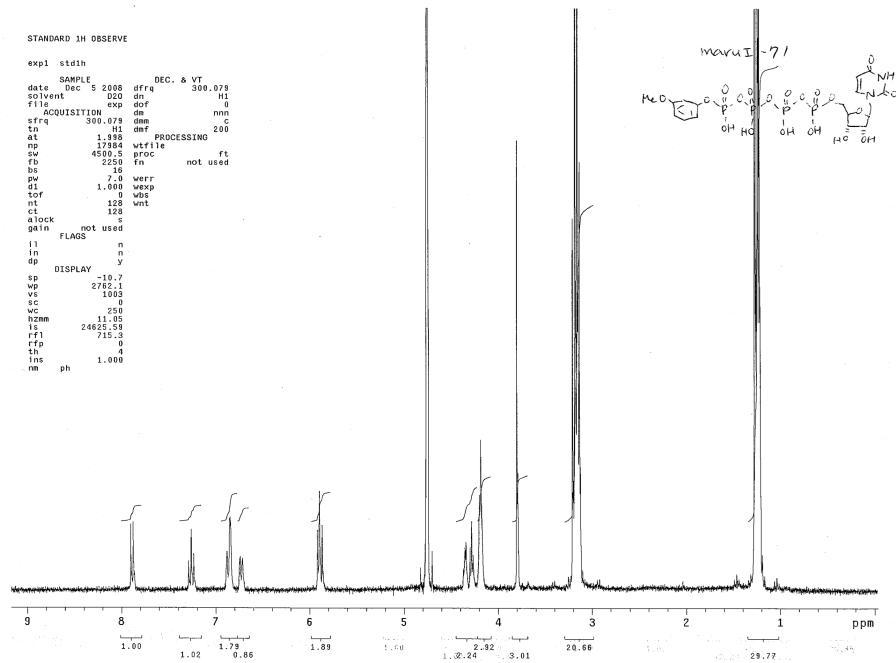


³¹P NMR

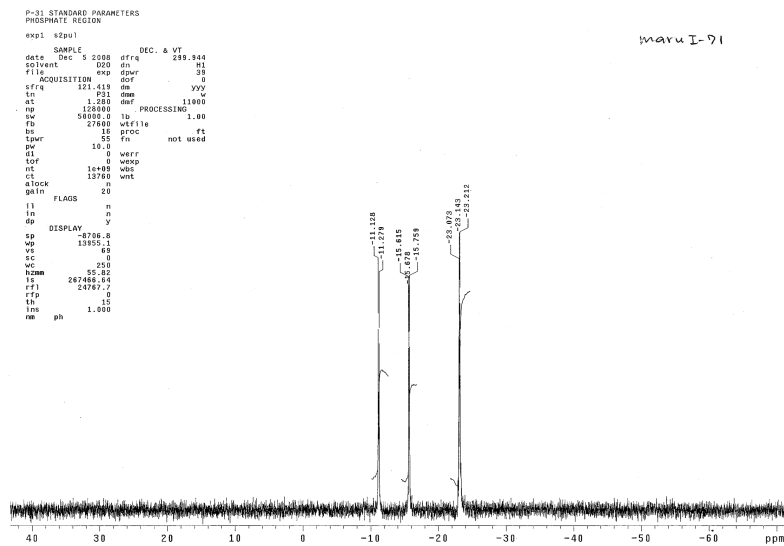


Uridine-5'-(3-methoxyphenyl)- tetraphosphate triethylammonium salt (23)

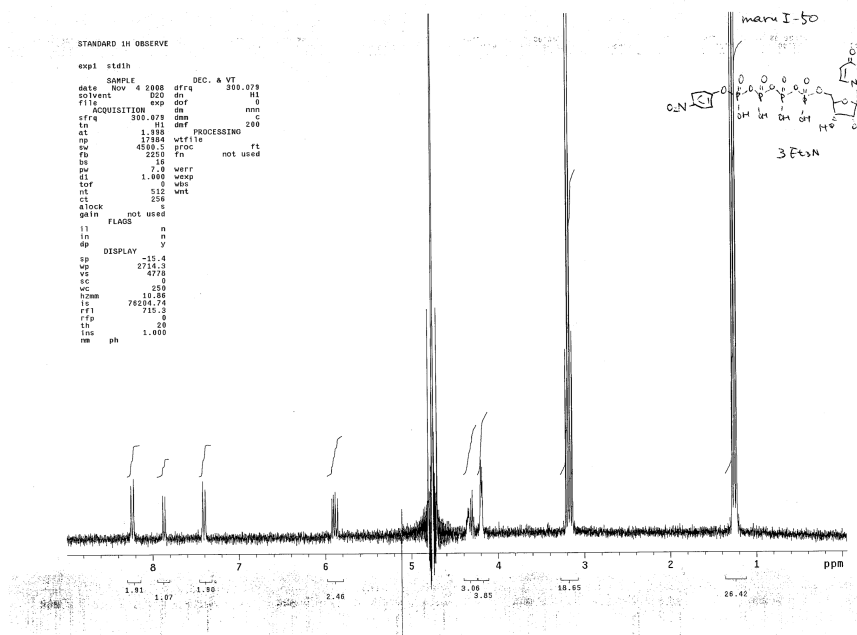
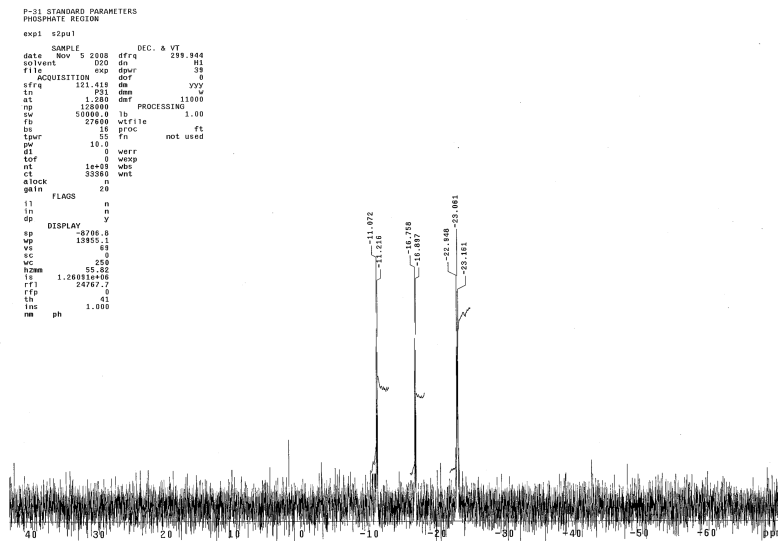
¹H NMR



³¹P NMR

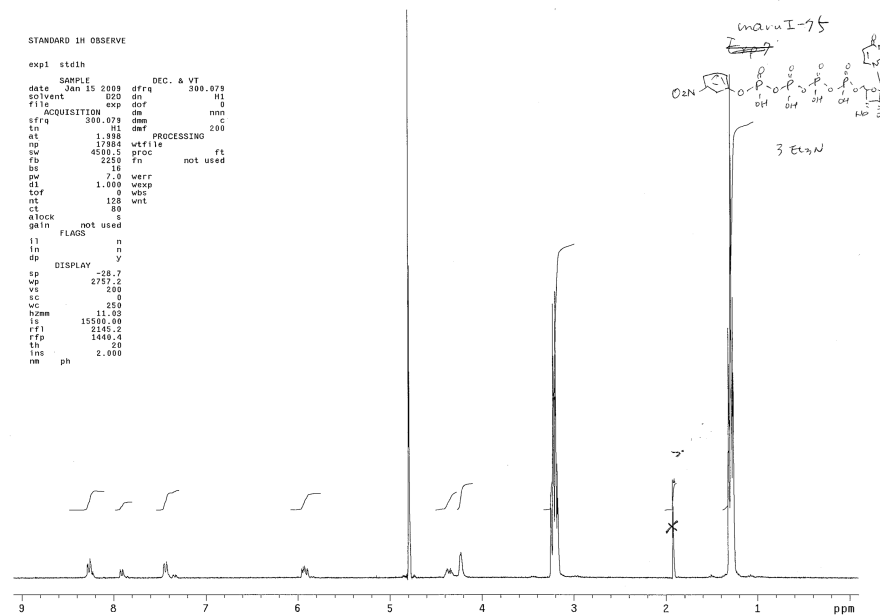


Uridine-5'-(4-nitrophenyl)- tetraphosphate triethylammonium salt (24)

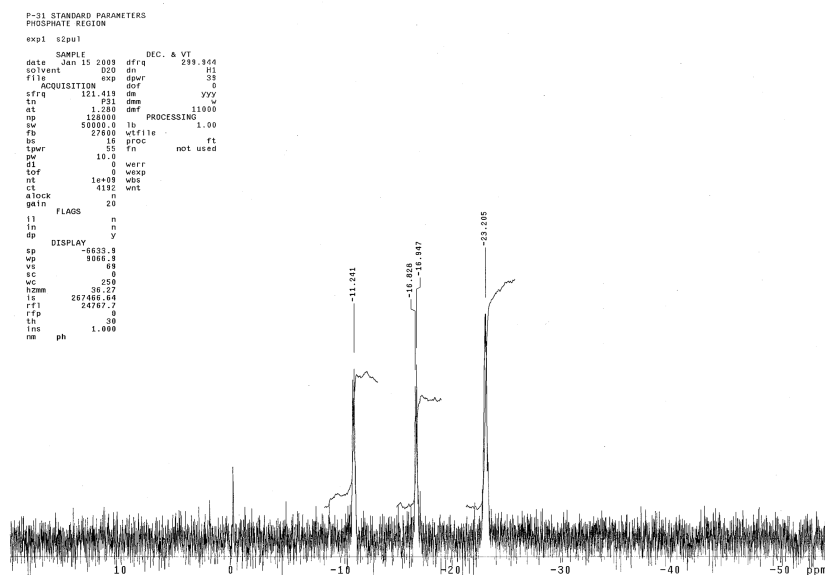
 ^1H NMR ^{31}P NMR

Uridine-5'-(3-nitrophenyl)- tetraphosphate triethylammonium salt (25)

^1H NMR

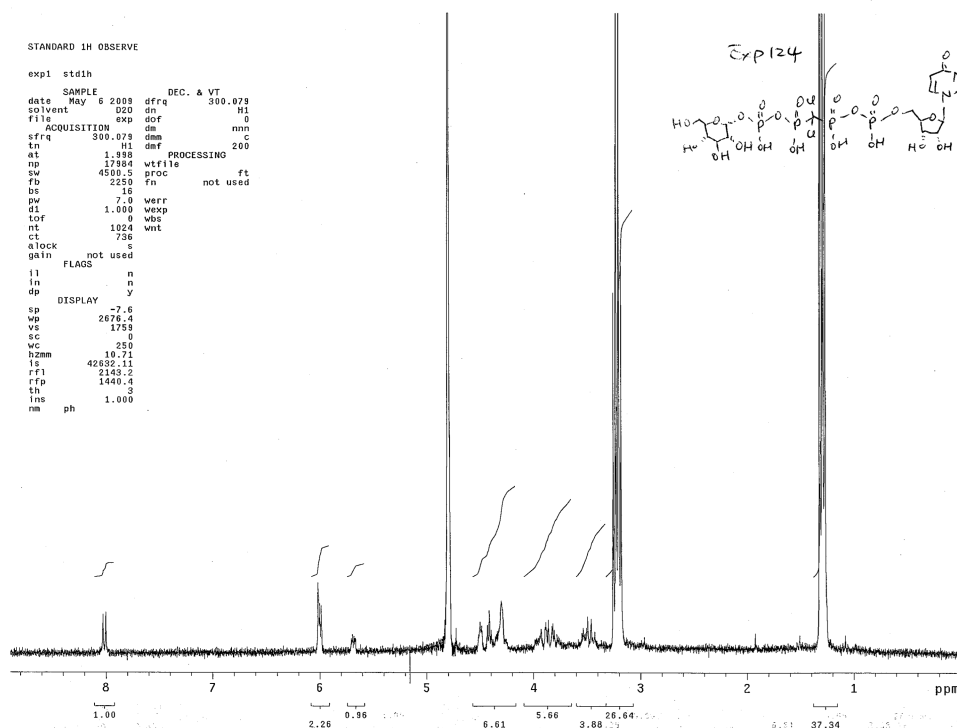


^{31}P NMR

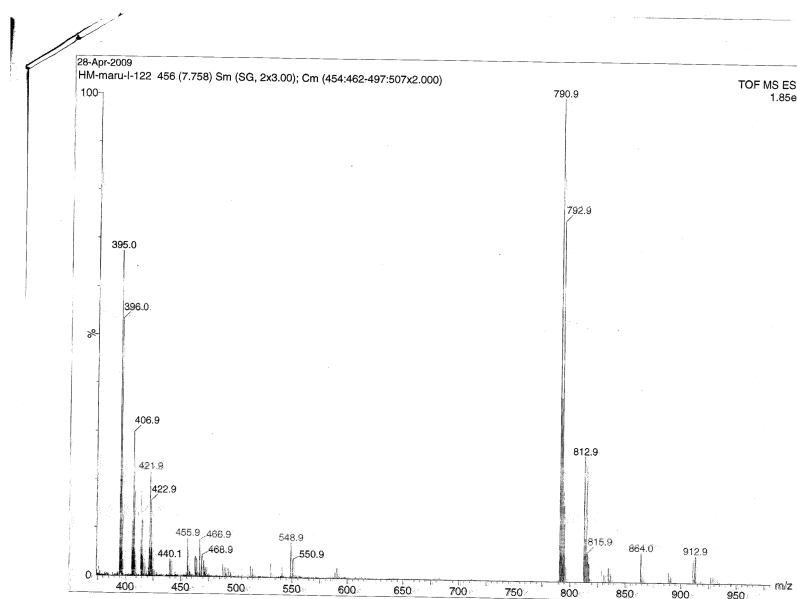


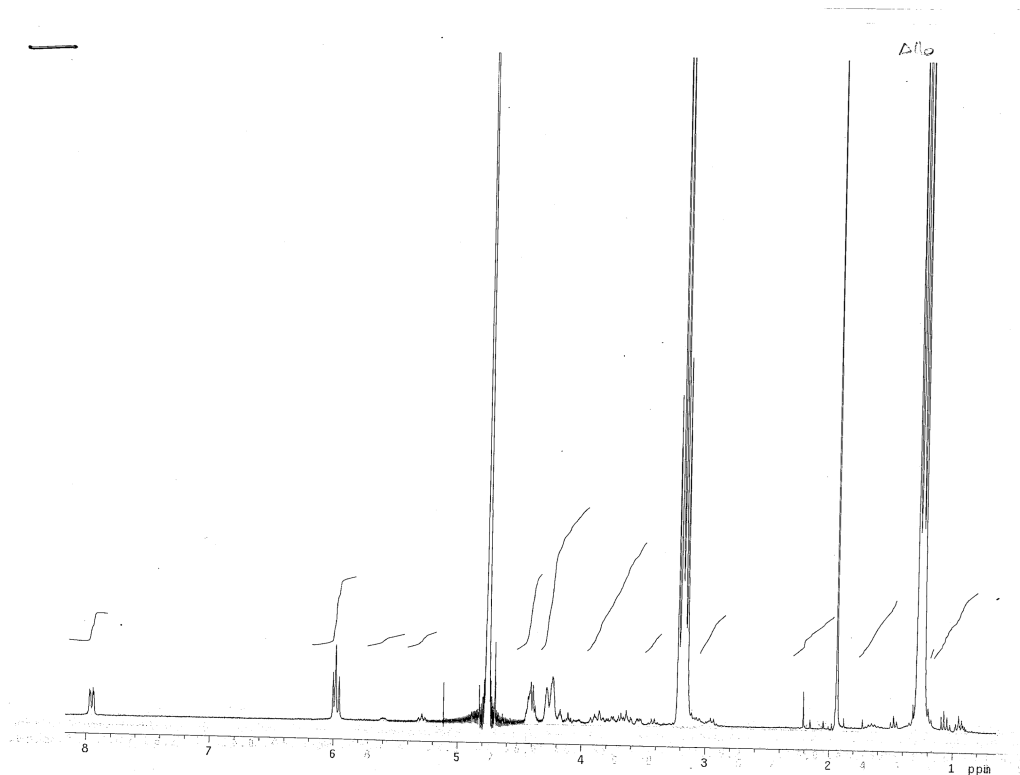
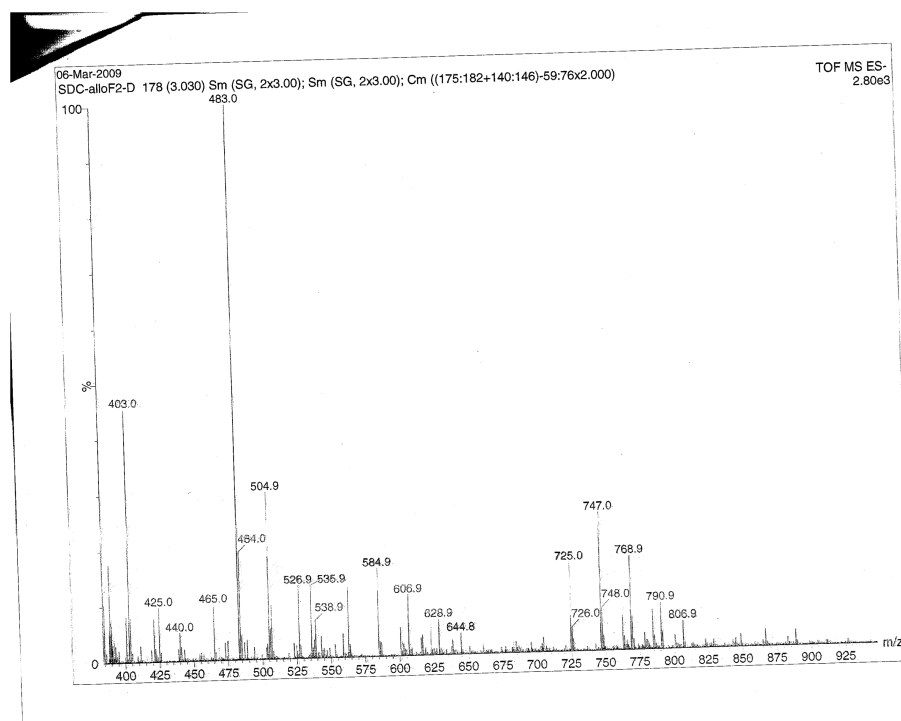
Uridine-5'-glucose-1'- β,γ -dichloromethylene tetraphosphate triethylammonium salt

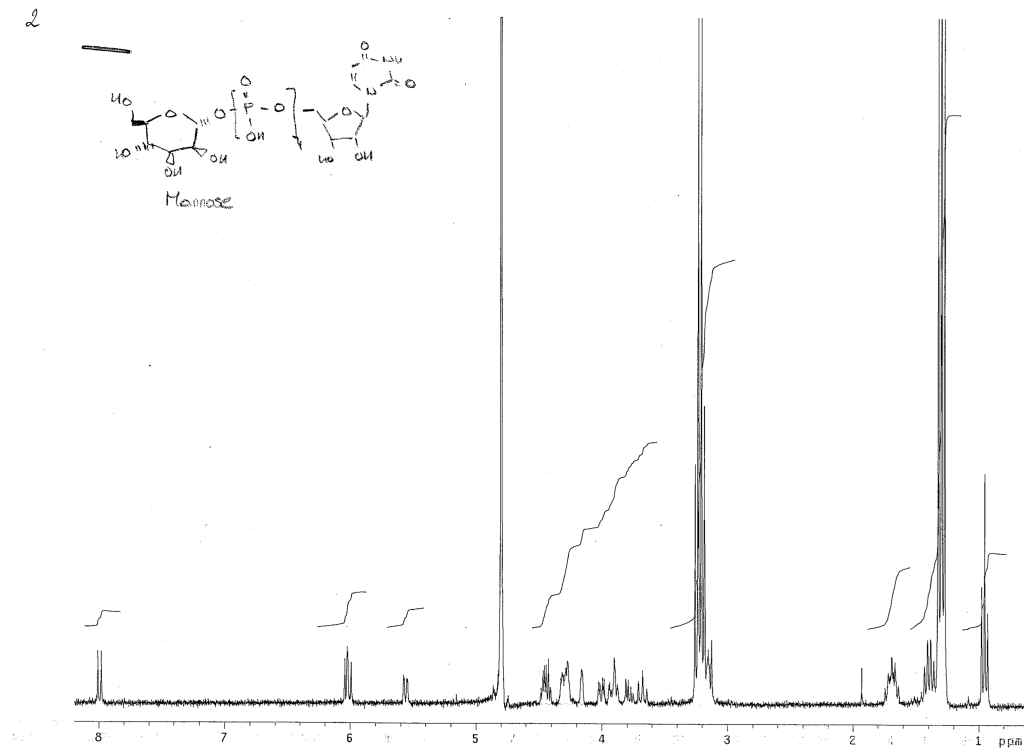
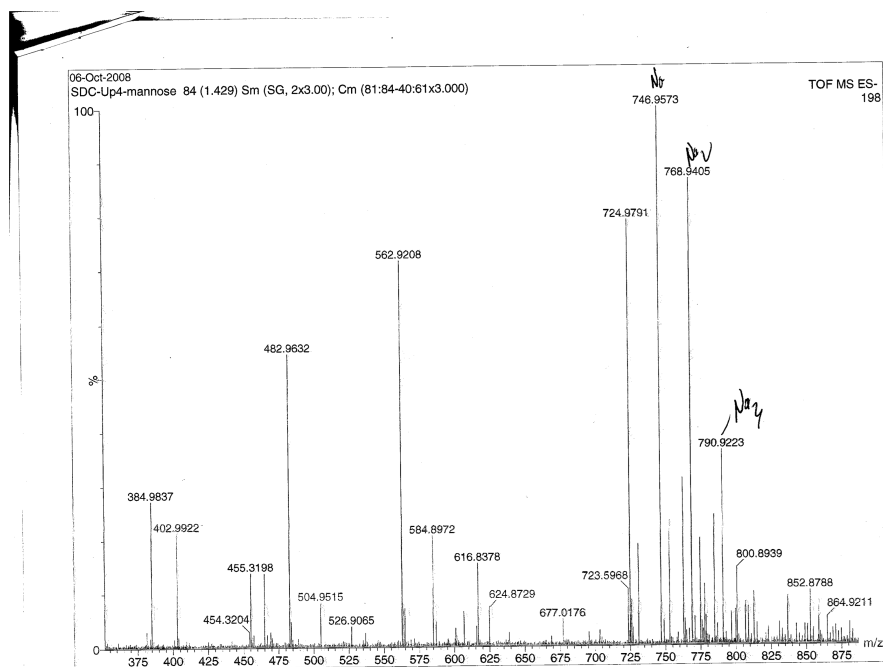
(26)

 ^1H NMR

Mass

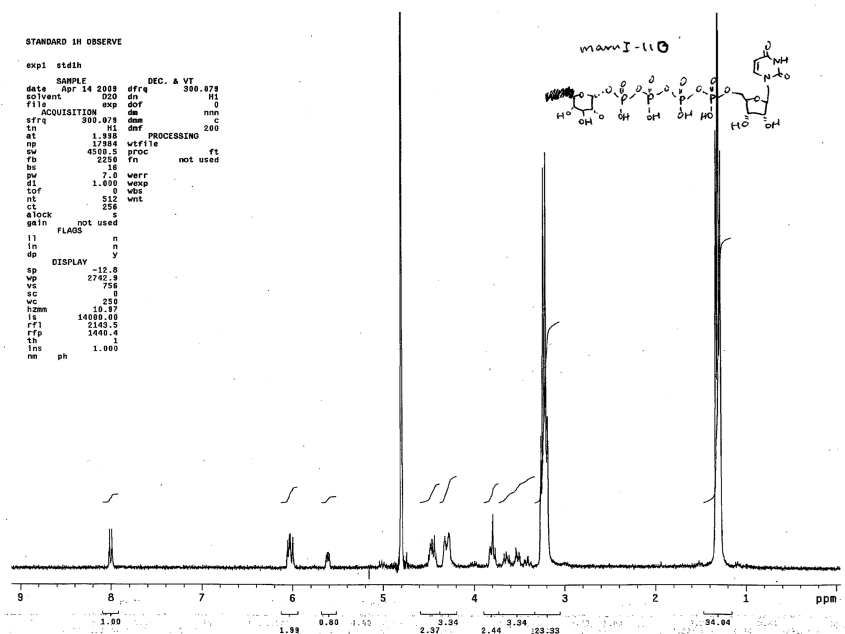


Uridine-5'-allose-1'-tetrphosphate triethylammonium salt (28)**¹H NMR****Mass**

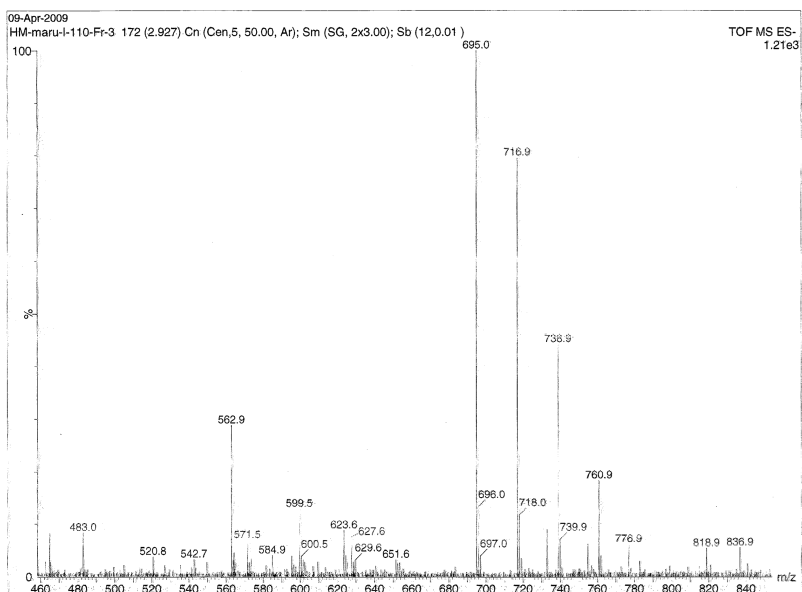
Uridine-5'-mannose-1'-tetraphosphate triethylammonium salt (29)**¹H NMR****Mass**

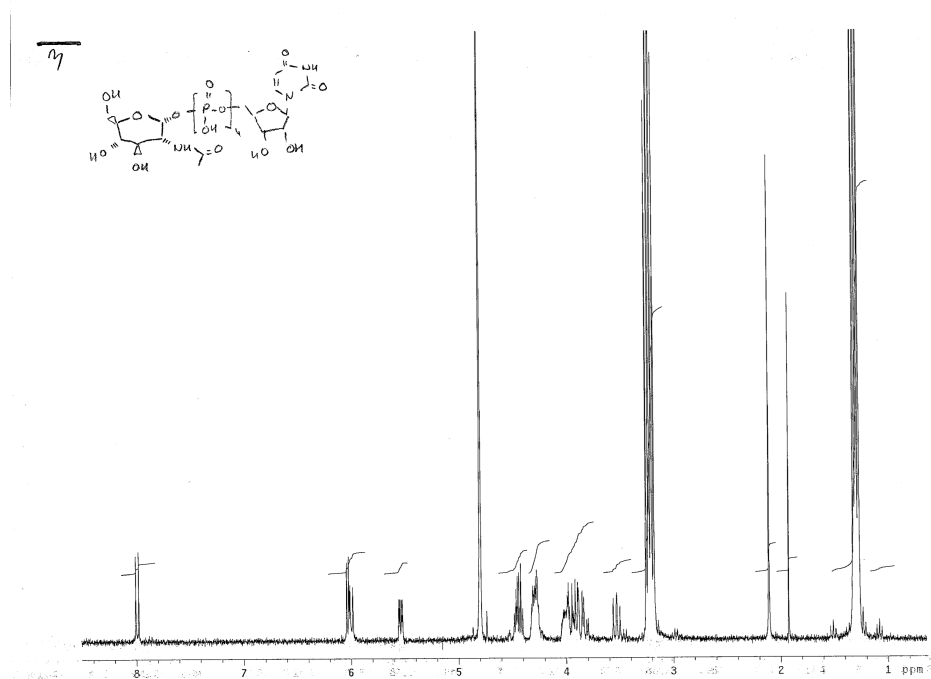
Uridine-5'-xylose-1'-tetraphosphate triethylammonium salt (30)

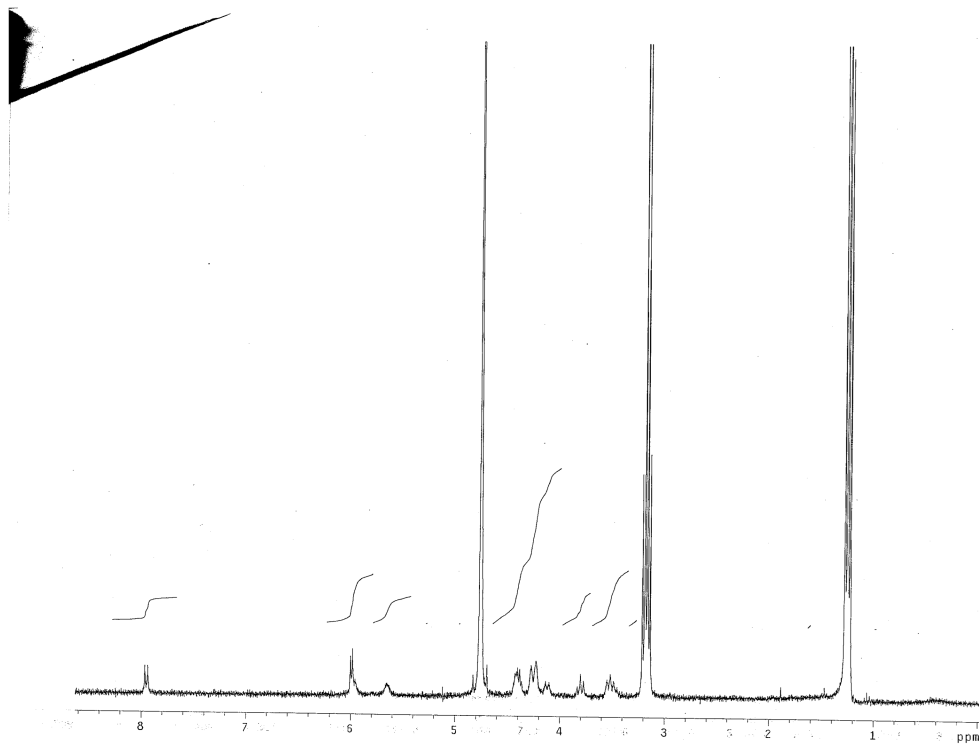
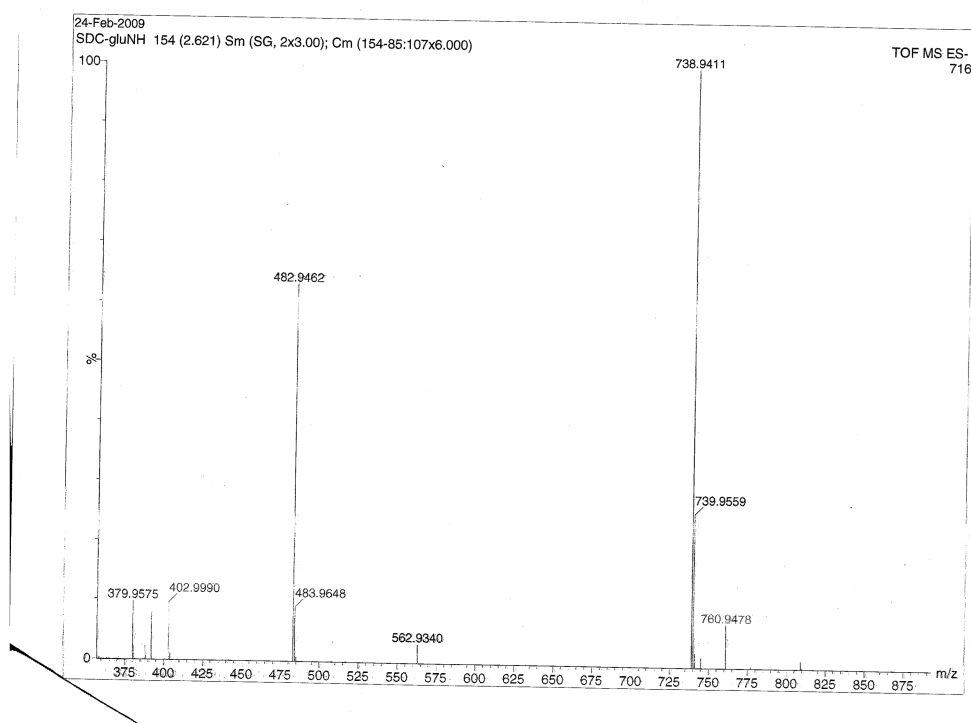
¹H NMR



Mass

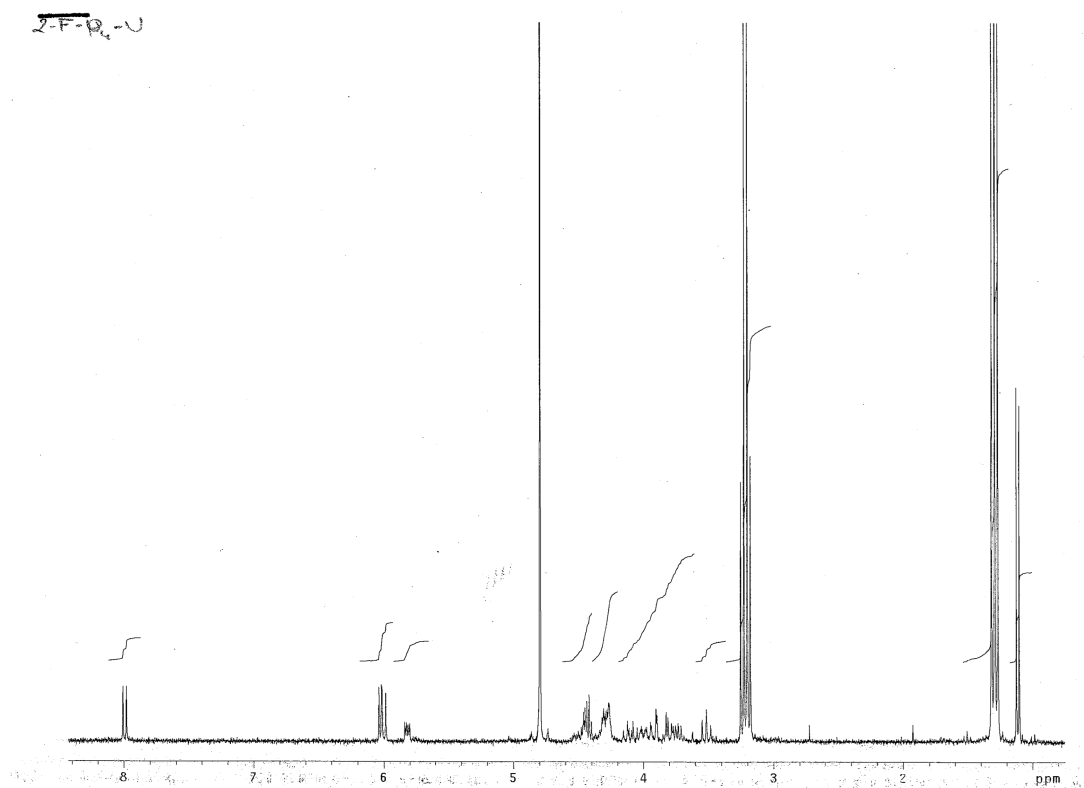


P¹-(Uridine-5'-)-P⁴-(2'-deoxy-2'-acetaminoglucose-1'-)tetraphosphate**triethylammonium salt (31)****¹H NMR****¹⁹F NMR** η

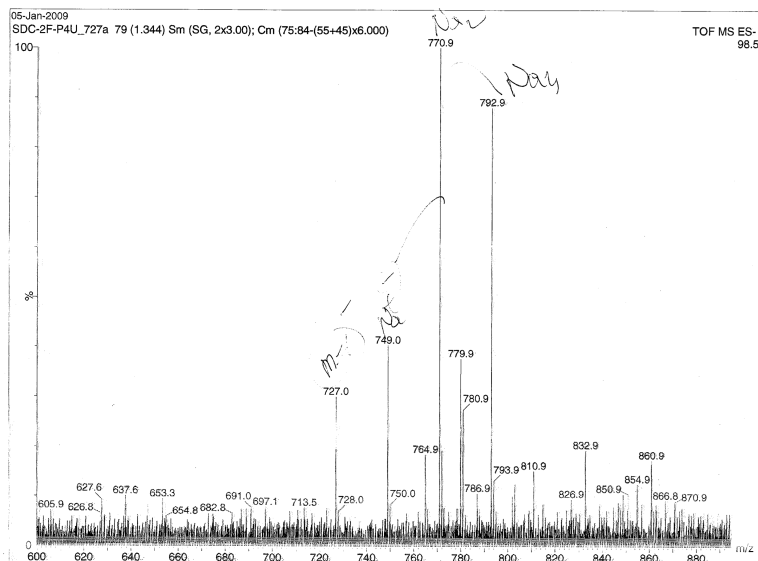
P¹-(Uridine-5'-)-P⁴-(glucuronic acid-1'-)tetraphosphate triethylammonium salt (32)**¹H NMR****Mass**

P¹-(Uridine-5'-)-P⁴-(2'-deoxy-2'-fluoroglucose-1'-)-tetrphosphate triethylammonium salt (33)

¹H NMR

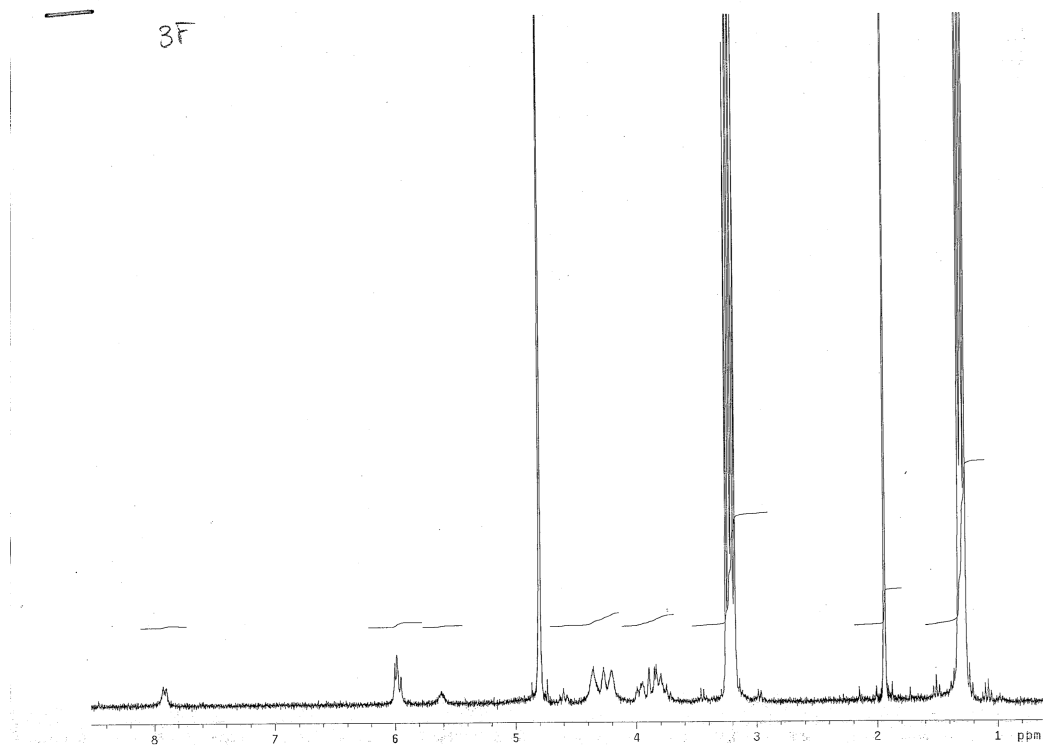


Mass

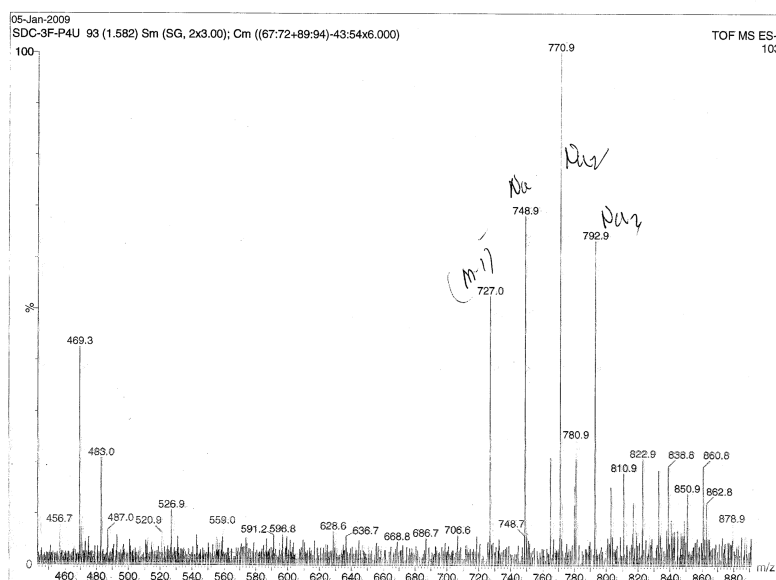


P¹-(Uridine-5'-)-P⁴-(3'-deoxy-3'-fluoroglucose-1'-)-tetrphosphate triethylammonium salt (34)

¹H NMR

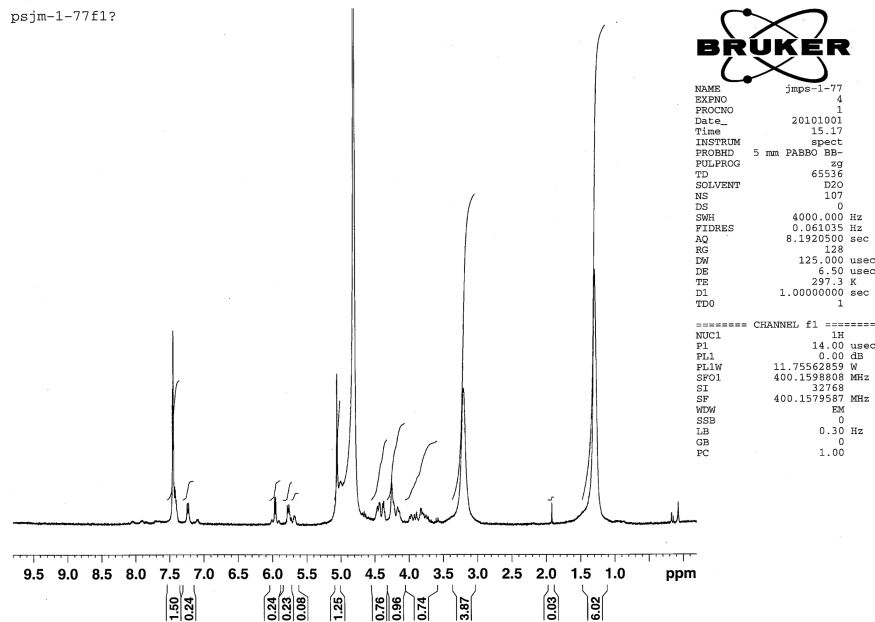


Mass

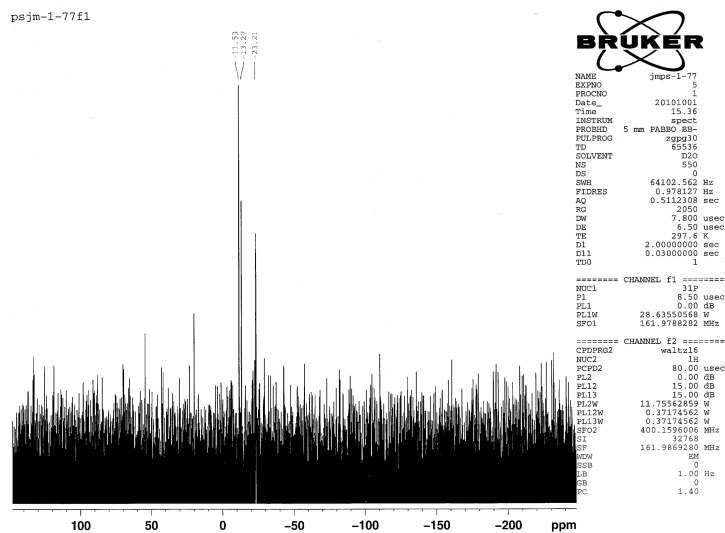


**P¹-(N⁴-Benzyloxycytidine-5')P⁴-(3'-deoxy-3'-fluoroglucose-1')tetrphosphate
triethylammonium salt (35)**

¹H NMR



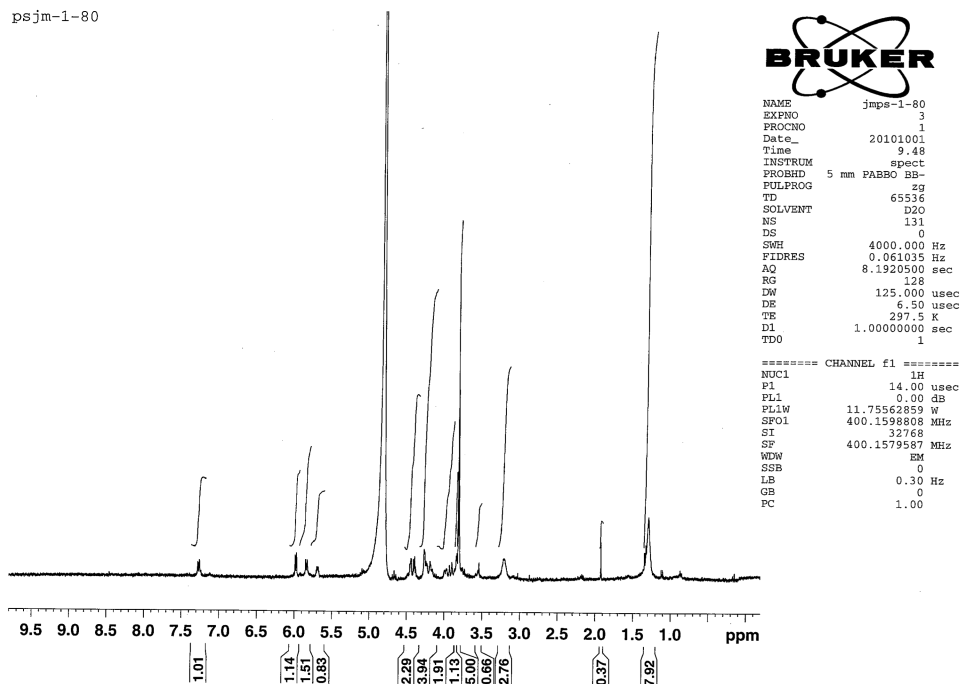
³¹P NMR



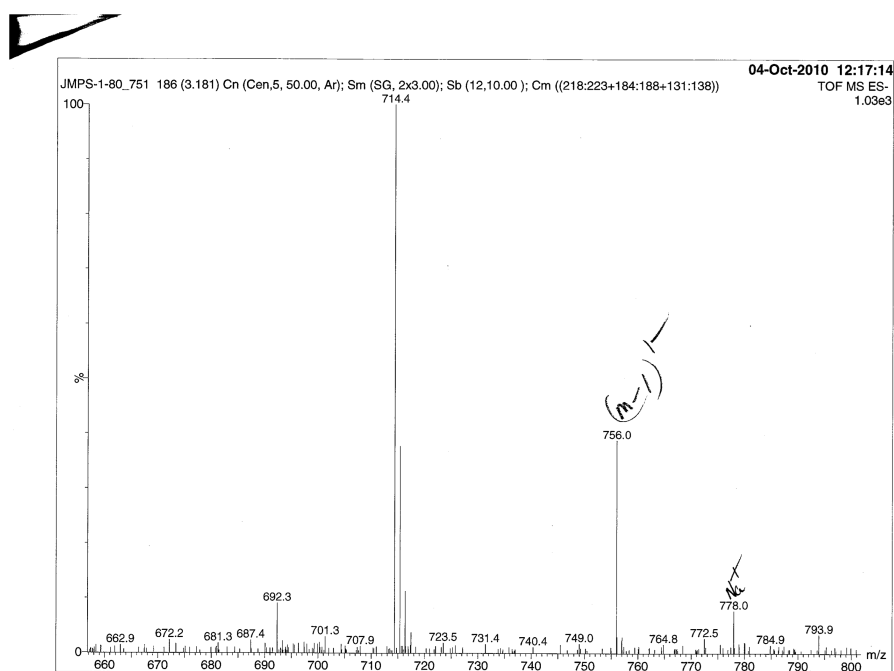
**P¹-(N⁴-Methoxycytidine-5'-)-P⁴-(3'-deoxy-3'-fluoroglucose-1'-)-tetrphosphate
triethylammonium salt (36)**

¹H NMR

psjm-1-80

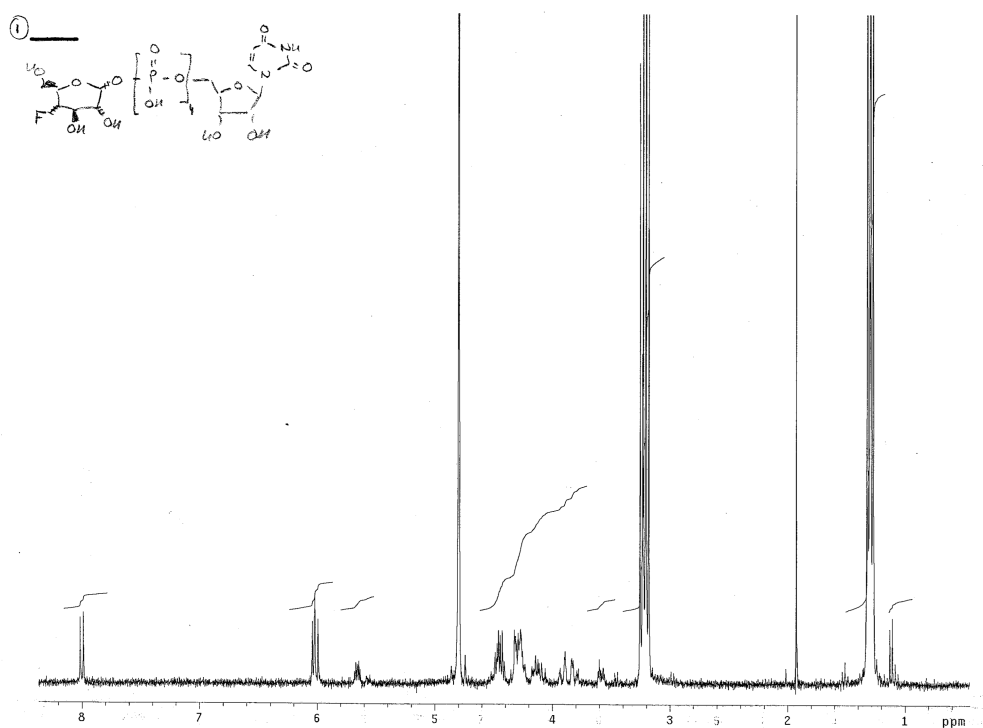


Mass



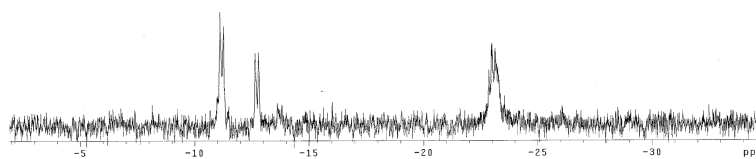
P¹-(Uridine-5'-)P⁴-(4'-deoxy-4'-fluoroglucose-1'-)tetraphosphate triethylammonium salt (37)

¹H NMR



¹⁹F NMR

4F



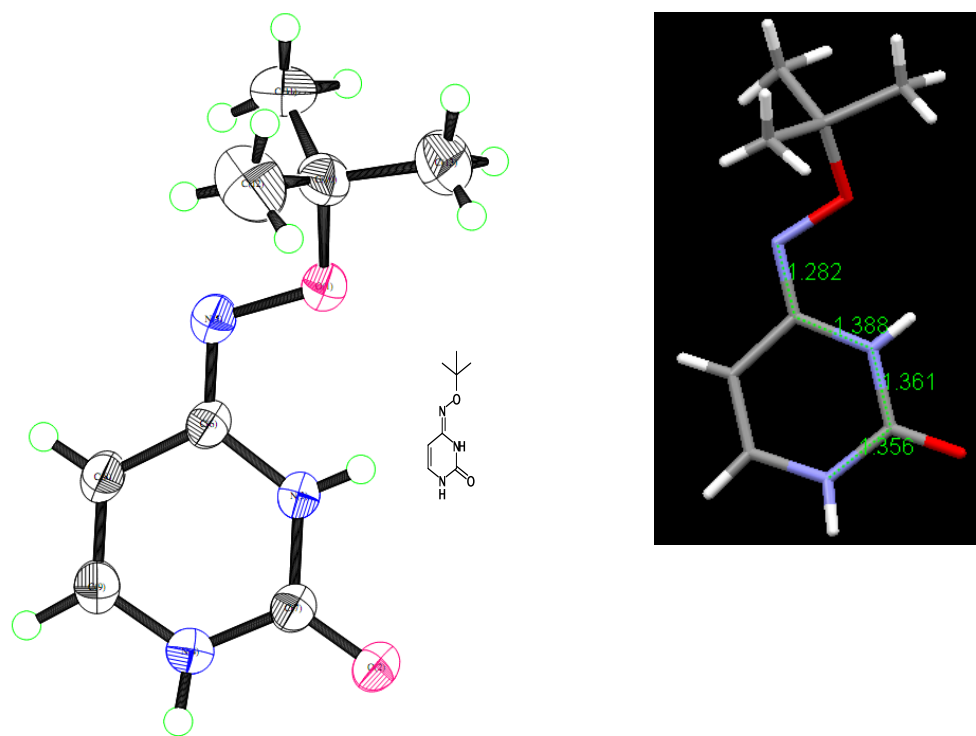


Figure 1. Structure of *N*⁴-*t*-BuO-cytosine determined by x-ray crystallography (bond lengths in Å).

Identification code	SMX10004
Empirical formula	C ₈ H ₁₃ N ₃ O ₂
Formula weight	183.21
Temperature	296.2 K
Wavelength	0.7107 Å
Crystal system	Monoclinic
Space group	C 2/c
Unit cell dimensions	a = 21.98(1) Å alpha = 90 deg. b = 7.543(5) Å beta = 105.16(2) deg. c = 12.120(6) Å gamma = 90 deg.
Volume	1939(2) Å ³
Z	8
Density (calculated)	1.255 Mg/m ³
Absorption coefficient	0.092 mm ⁻¹
F(000)	784
Crystal size	0.60 x 0.60 x 0.40 mm
Theta range for data collection	3.22 to 27.48 deg.
Index ranges	-28 ≤ h ≤ 28, -9 ≤ k ≤ 9, -15 ≤ l ≤ 15
Reflections collected	9631
Independent reflections	2221 [R(int) = 0.01460]
Refinement method	Full-matrix least-squares on F ²
Data / restraints / parameters	2221 / 0 / 118
Goodness-of-fit on F ²	1.081
Final R indices [I > 2σ(I)]	R1 = 0.0534, wR2 = 0.1651
R indices (all data)	R1 = 0.0584, wR2 = 0.1707
Largest diff. peak and hole	0.378 and -0.259 e.Å ⁻³

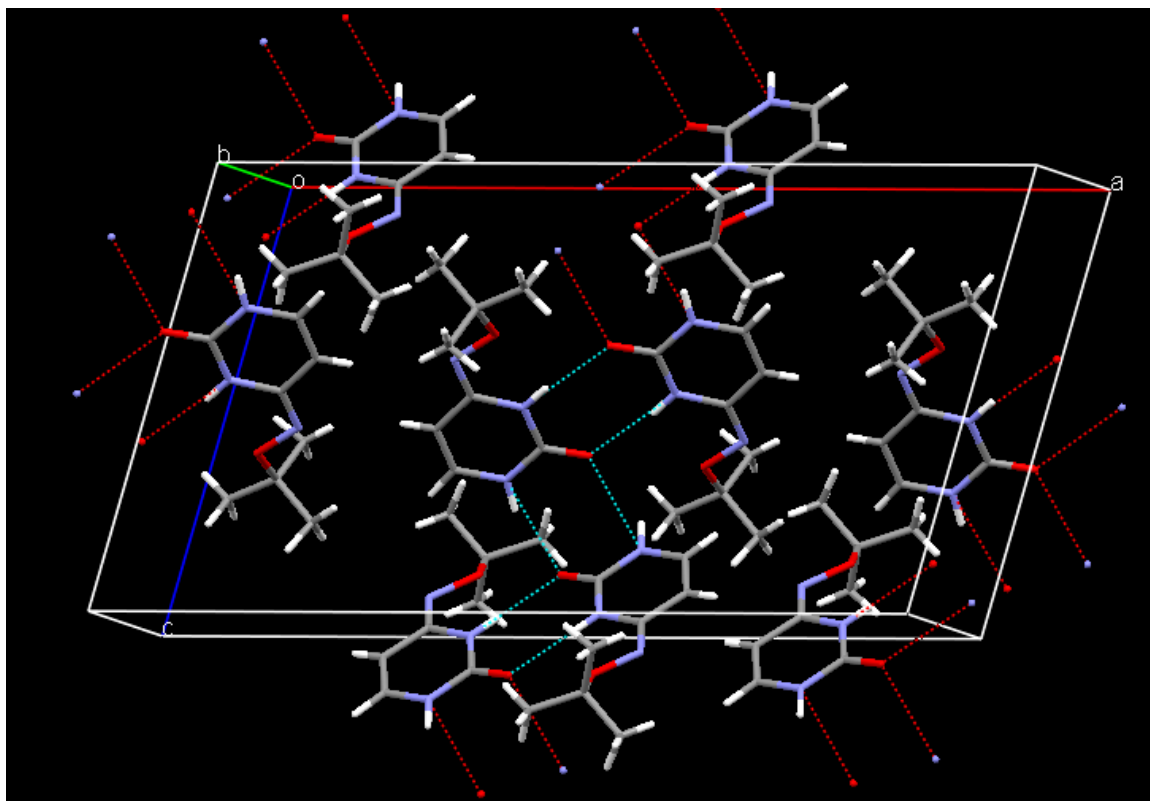


Figure 2. Crystal lattice of *N*⁴-*t*-BuO-cytosine determined by x-ray crystallography.

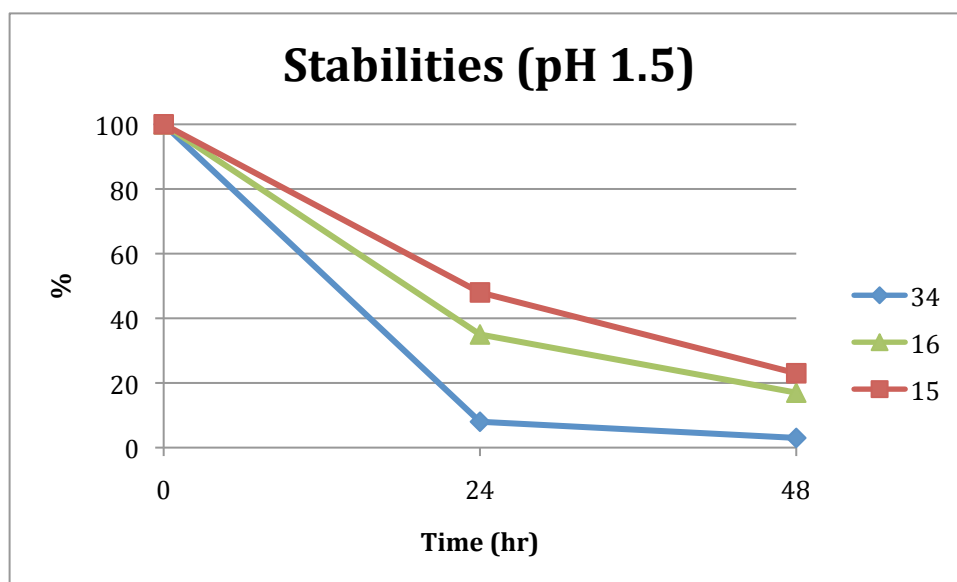
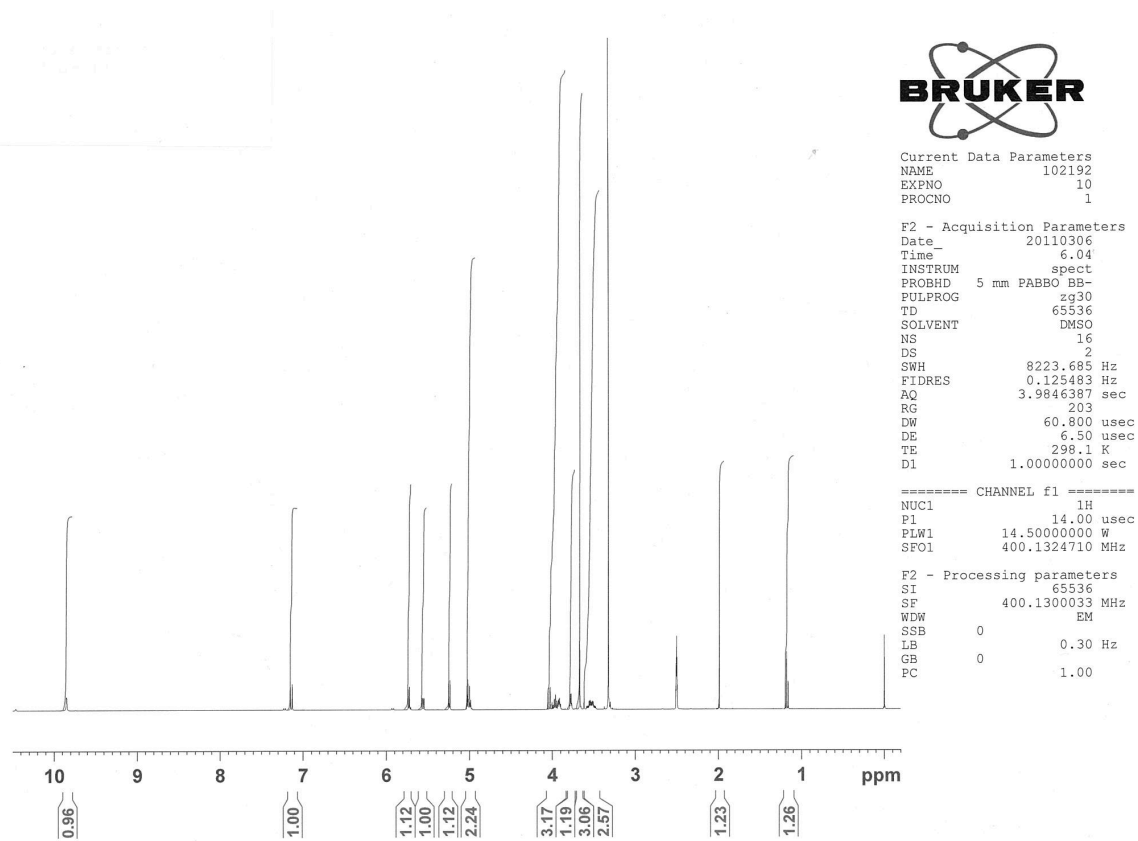
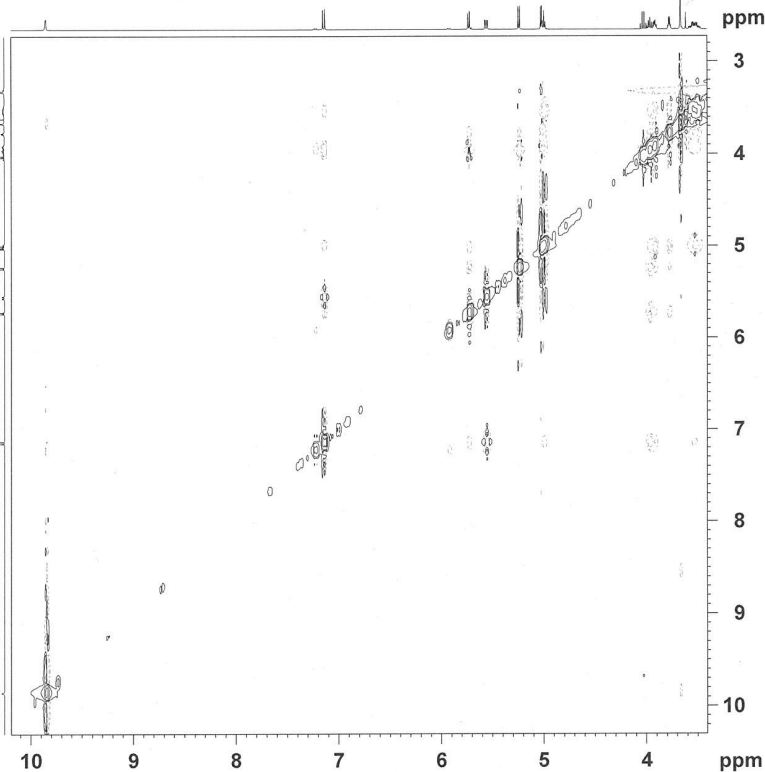


Figure 3. Stability data at pH 1.5 (37°C) for compounds **15**, **16**, and **34**.

Figure 4. NOESY experiment for compound 39





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 PROCNO 1

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 TD 2048
 SOLVENT DMSO
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 SWH 4545.455 Hz
 FIDRES 2.219460 Hz
 AQ 0.2253300 sec
 RG 114
 DW 110.000 usec
 DE 6.50 usec
 TE 298.1 K
 D0 0.0009217 sec
 D1 2.02339492 sec
 D8 0.30000001 sec
 D11 0.03000000 sec
 D12 0.00020000 sec
 D16 0.00020000 sec
 INO 0.00022000 sec

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 P2 28.00 usec
 P17 2500.00 usec
 PLW1 14.50000000 W
 PLW0 3.15779996 W
 SFO1 400.131993 MHz

----- GRADIENT CHANNEL -----
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 GPZ1 40.00 %
 P16 1000.00 usec

F1 - Acquisition parameters
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 FIDRES 17.755682 Hz
 SW 11.360 ppm
 FMODE States-TPPI

F2 - Processing parameters
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 WW QFINE
 SSB 2
 LB 0 Hz
 GB 0
 PC 1.00

F1 - Processing parameters
 SI 1024
 MC2 States-TPPI
 SF 400.1300033 MHz
 WW QFINE
 SSB 2
 LB 0 Hz
 GB 0

