

Supporting Information

Structure-Activity Relationship of Truncated 2,8-Disubstituted-adenosine Derivatives as Dual A_{2A}/A₃ Adenosine Receptor Antagonists and their Cancer Immunotherapeutic Activity

Gibae Kim,¹ Xiyan Hou,^{1,2} Woong Sub Byun,^{1,3} Gyudong Kim,^{1,4} Dnyandev B. Jarhad,¹ Grim Lee,¹ Young Eum Hyun,¹ Jinha Yu,^{1,5} Chang Soo Lee,¹ Shuhao Qu,¹ Eugene Warnick,⁶ Zhan-Guo Gao,⁶ Ji Yong Kim,⁷ Seunghee Ji,⁸ Hyunwoo Shin,⁸ Jong-Ryoul Choi,⁸ Kenneth A. Jacobson,⁶ Hyuk Woo Lee,⁷ Sang Kook Lee,¹ Lak Shin Jeong^{1,7,*}

¹Research Institute of Pharmaceutical Sciences, College of Pharmacy, Seoul National University, Seoul 08826, Republic of Korea, ²College of Life Science, Dalian Minzu University, Dalian 116600, People's Republic of China, ³Department of Chemical and Systems Biology, Chem-H and Stanford Cancer Institute, Stanford School of Medicine, Stanford University, Stanford, California 94305, United States, ⁴College of Pharmacy & Research Institute of Drug Development, Chonnam National University, Gwangju 61186, Republic of Korea, ⁵College of Pharmacy and Graduate School of Pharmaceutical Sciences, Ewha Womans University, Seoul 03760, South Korea, ⁶Molecular Recognition Section, Laboratory of Bioorganic Chemistry, National Institute of Diabetes, and Digestive and Kidney Disease, National Institutes of Health, Bethesda, Maryland 20892, USA, and ⁷Future Medicine Co., Ltd, Seoul 06665, Republic of Korea, ⁸HK inno.N Corp., Seoul 04551, Republic of Korea.

*Corresponding authors: lakjeong@snu.ac.k

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Additional *in vitro* data and procedures

Table S1. The cAMP functional assay of A_{2A}AR with 100 nM of full agonist CGS21680

Compound	% inhibition, A _{2A} AR cAMP functional assay
5a	53.47%
5d	63.82%
5g	12.31%

Table S2. The result of *in vitro* CYP inhibition assay

Compound	% inhibition at 10 μ M				
	CYP1A2	CYP2C9	CYP2C19	CYP2D6	CYP3A4
5d	14%	4%	6%	3%	0%

Table S3. The result of *in vitro* hERG inhibition assay

Compound	% inhibition		
	1 μ M	3 μ M	10 μ M
5d	13%	18%	31%

Cyclic AMP Accumulation Assay with CGS-21680. Human adenosine receptor functional experiments were carried out in A_{2A}AR-HEK293 cell line for hA_{2A}AR. The cells are seeded on the 96 well culture plate (NUNC 136101) with stimulation buffer (1X HBSS, Gibco 14025092; 7.5% BSA stabilizer; 250 mM IBMX, Sigma I5879; 1 M HEPES). 10 μ M of test compound and 100 nM of CGS-21680 (**26**) are added, then the cells are centrifuged at 500 rpm for 3 min, followed by incubated at 30 °C for 30 min. After incubation, the amount of cAMP is determined using LANCE Ultra cAMP detection kit (Perkin Elmer TRF0262).

***In vitro* CYP inhibition assay.** Human liver microsomes (0.25 mg/mL), 0.1 M phosphate buffer (pH 7.4), a cocktail of five coenzyme substrates (Phenacetin 50 μ M, Diclofenac 10 μ M, S-mephenytoin 100 μ M, Dextromethorphan 5 μ M, Midazolam 2.5 μ M), and test compound (10 μ M

concentration) was pre-incubated for 5 min at 37 °C. NADPH generation system solution was added and incubated for 15 min at 37 °C. In order to terminate the reaction, acetonitrile solution containing an internal standard (Terfenadine) was added and centrifuged for 5 min (14,000× g rpm, 4 °C). The supernatant was injected into the LC-MS/MS system to analyze the metabolites of the substrates simultaneously. Metabolites of each substrate produced during the reaction were analyzed using the Shimadzu Nexera XR system and TSQ vantage (Thermo). Kinetex C18 column (2.1 × 100 mm, 2.6 μm particle size; Phenomenex, USA) was used for HPLC. The mobile phase used contained 0.1% formic acid in distilled water (A) and 0.1% formic acid containing acetonitrile (B). The generated metabolites were quantified using MRM (Multiple Reaction Monitoring) and Xcalibur (version 1.6.1) was used for data analysis.^{1,2}

In vitro hERG assay. CHO cells stably expressing hERG potassium channels from Sophion Biosciences were used for this test. The cells were cultured in a humidified and air-controlled (5% CO₂) incubator at 37 °C. The CHO cells which were at least two days after plating and more than 75% confluent would be used for experiments. Before testing, cells were harvested using TrypLE and resuspended in the physiological solution at the room temperature. The physiological solution and external solution were prepared at least one month. The intracellular solution was prepared in batches aliquoted, and stored at 4 °C until used. Test compounds were dissolved in 100% DMSO at 10 mM to obtain stock solutions for different test concentrations. Then the stock solutions were further diluted into external solution to achieve final concentrations for testing. Visual check for precipitation was conducted before testing. Final DMSO concentration in extracellular solution was not more than 0.30% for the test compounds. hERG SyncroPatch(384PE) assay was conducted at room temperature. The Setup, Prime Chip, Catch and Seal Cells, Amplifier Settings, Voltage and Application Protocols were established with Biomek Software (Nanion). One addition of 40 μL of the vehicle was applied, followed by 300s for a baseline period. Then the doses of the compounds were added with 40 μL. The exposure of test compound at each concentration was no less than 300s. The recording for the whole process had to pass the quality control or the well was abandoned and the compound was retested, all automatically set by PatchControl. Three concentrations (1.00 μM, 3.00 μM, and 10.00 μM) were tested for each compound. Minimum 2 replicates per concentration were obtained.

Single-Crystal X-ray crystallography data of compounds **11b**, **22a** and **22b**.

Compound **11b**, **22a** or **22b** (~10 mg) was dissolved in ethyl acetate and dichloromethane (1.0 mL, 1:4) in a glass vial. After staying at room temperature, the single crystals of **11b**, **22a** or **22b** were obtained. X-ray diffraction data of **11b** (CCDC No: 2164196), **22a** (CCDC No: 2164194), and **22b** (CCDC No: 2258518) were obtained on a SuperNova, Dual, Cu at zero, AtlasS2 diffractometer. Using Olex2³, the structure was solved with the ShelXT⁴ structure solution program using Intrinsic Phasing and refined with the ShelXL⁴ refinement package using Least Squares minimization.

Single-Crystal X-ray crystallography of compound **11b**

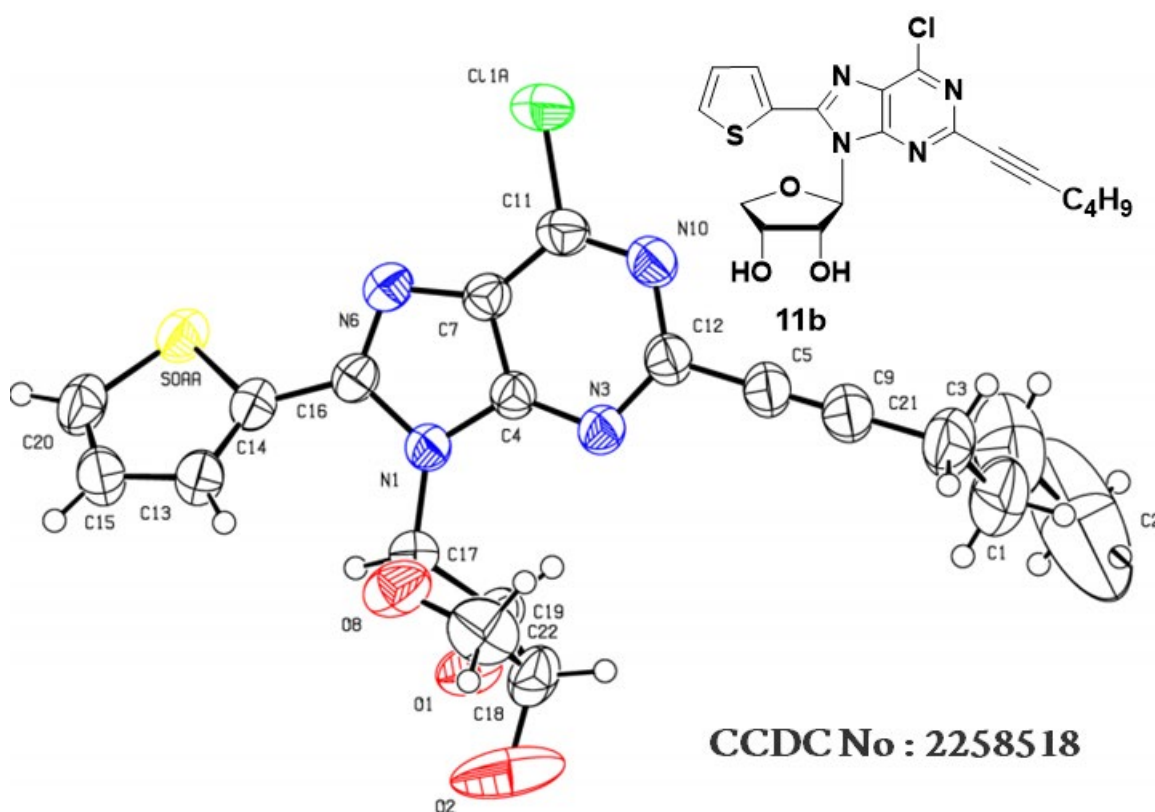


Figure S1. ORTEP diagram of compound **11b** showing thermal ellipsoid at 50% probability.

Table S4. Crystal data and structure refinement for **11b**

Empirical formula	C ₁₉ H ₁₉ ClN ₄ O ₃ S
Formula weight	418.89
Temperature/K	293(1)
Crystal system	monoclinic
Space group	C2
a/Å	25.4206(7)
b/Å	7.3860(2)
c/Å	10.6375(3)
α/°	90
β/°	96.378(2)
γ/°	90
Volume/Å ³	1984.90(10)
Z	4
ρ _{calc} /cm ³	1.402
μ/mm ⁻¹	2.929
F(000)	872.0
Crystal size/mm ³	0.563 × 0.19 × 0.087
Radiation	CuKα (λ = 1.54184)
2θ range for data collection/°	6.998 to 147.472
Index ranges	-31 ≤ h ≤ 26, -8 ≤ k ≤ 8, -13 ≤ l ≤ 12
Reflections collected	3649
Independent reflections	2927 [R _{int} = 0.0113, R _{sigma} = 0.0187]
Data/restraints/parameters	2927/1/256
Goodness-of-fit on F ²	1.036

Table S5. Crystal data and structure refinement for **22a**

Empirical formula	C ₁₉ H ₁₉ ClN ₄ O ₃ S
Formula weight	418.89
Temperature/K	293(2)
Crystal system	orthorhombic
Space group	P2 ₁ 2 ₁ 2 ₁
a/Å	7.1613(18)
b/Å	10.7229(15)
c/Å	25.009(4)
α/°	90
β/°	90
γ/°	90
Volume/Å ³	1920.5(6)
Z	4
ρ _{calc} /cm ³	1.449
μ/mm ⁻¹	0.337
F(000)	872.0
Crystal size/mm ³	0.262 × 0.114 × 0.075
Radiation	MoKα (λ = 0.71073)
2θ range for data collection/°	5.004 to 58.788
Index ranges	-8 ≤ h ≤ 8, -13 ≤ k ≤ 14, -25 ≤ l ≤ 34
Reflections collected	11286
Independent reflections	4489 [R _{int} = 0.0733, R _{sigma} = 0.1095]

Table S6. Crystal data and structure refinement for **22b**

Empirical formula	C ₁₉ H ₁₉ ClN ₄ O ₂ S ₂
Formula weight	434.97
Temperature/K	111.0(8)
Crystal system	orthorhombic
Space group	P2 ₁ 2 ₁ 2 ₁
a/Å	7.0735(7)
b/Å	24.677(4)
c/Å	12.1455(13)
α/°	90
β/°	90
γ/°	90
Volume/Å ³	2120.0(4)
Z	4
ρ _{calc} /cm ³	1.3627
μ/mm ⁻¹	0.399
F(000)	905.9
Crystal size/mm ³	0.466 × 0.394 × 0.229
Radiation	Mo Kα (λ = 0.71073)
2θ range for data collection/°	4.7 to 59.38
Index ranges	-8 ≤ h ≤ 9, -34 ≤ k ≤ 29, -15 ≤ l ≤ 16

Reflections collected	23486
Independent reflections	5357 [$R_{\text{int}} = 0.0955$, $R_{\text{sigma}} = 0.0926$]
Data/restraints/parameters	5357/12/255
Goodness-of-fit on F^2	1.038
Final R indexes [$I > 2\sigma(I)$]	$R_1 = 0.0902$, $wR_2 = 0.2143$
Final R indexes [all data]	$R_1 = 0.1161$, $wR_2 = 0.2353$
Largest diff. peak/hole / $e \text{ \AA}^{-3}$	1.00/-0.90

Reference

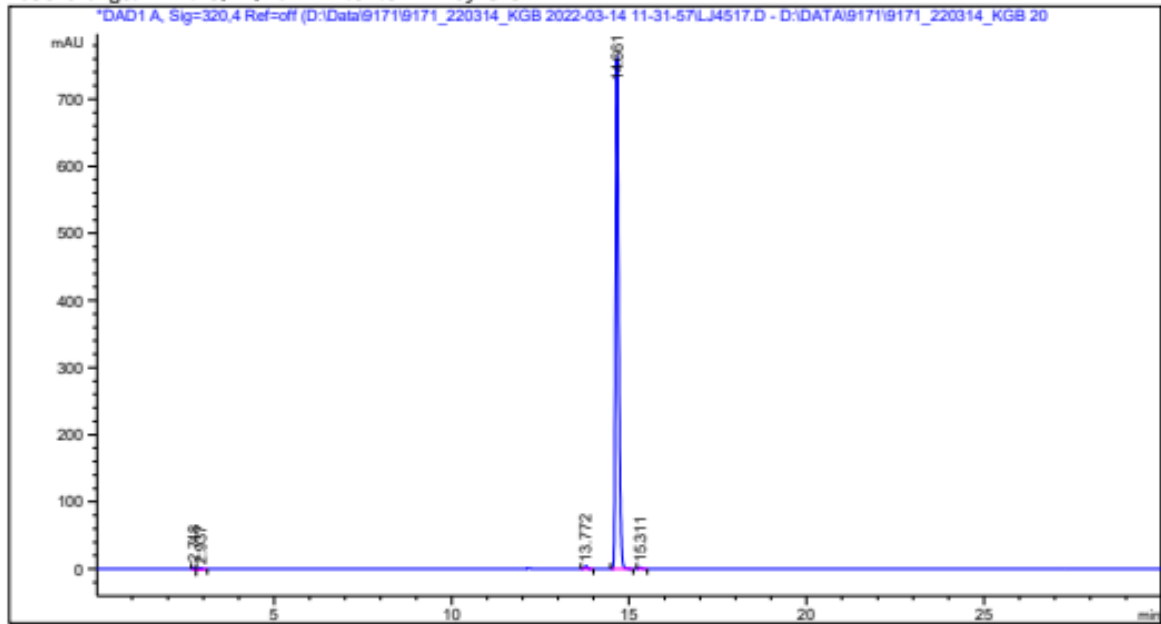
- (1) Kim, M.-J.; Kim, H.; Cha, I.-J.; Park, J.-S.; Shon, J.-H.; Liu, K.-H.; Shin, J.-G. High-throughput screening of inhibitory potential of nine cytochrome P450 enzymes *in vitro* using liquid chromatography/tandem mass spectrometry. *Rapid Commun. Mass Spectrom.* **2005**, *19*, 2651–2658.
- (2) Kerns, E.H.; Di, L. (Eds.) *Chapter 32—CYP Inhibition Methods. Drug-like Properties: Concepts, Structure Design and Methods*, Academic Press: San Diego, CA, USA, 2008, pp. 360–3
- (3) Dolomanov, O. V.; Bourhis, L. J.; Gildea, R. J.; Howard, J. A. K.; Puschmann, H. J. *Appl. Cryst.* **2009**, *42*, 339-341..
- (4) Sheldrick, G. M. *Acta Cryst.* **2015**, *A71*, 3-8.
- (5) Sheldrick, G. M. *Acta Cryst.* **2015**, *C71*, 3-8.

HPLC data of the representative compounds

*HPLC data of the compound **5d***

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Sample Name: LJ4517

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                                           Inj Volume: 10.000 µl
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Method)
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Area Percent Report
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Sorted By : Signal
Multiplier : 1.0000
Dilution : 1.0000
Do not use Multiplier & Dilution Factor with ISTDs

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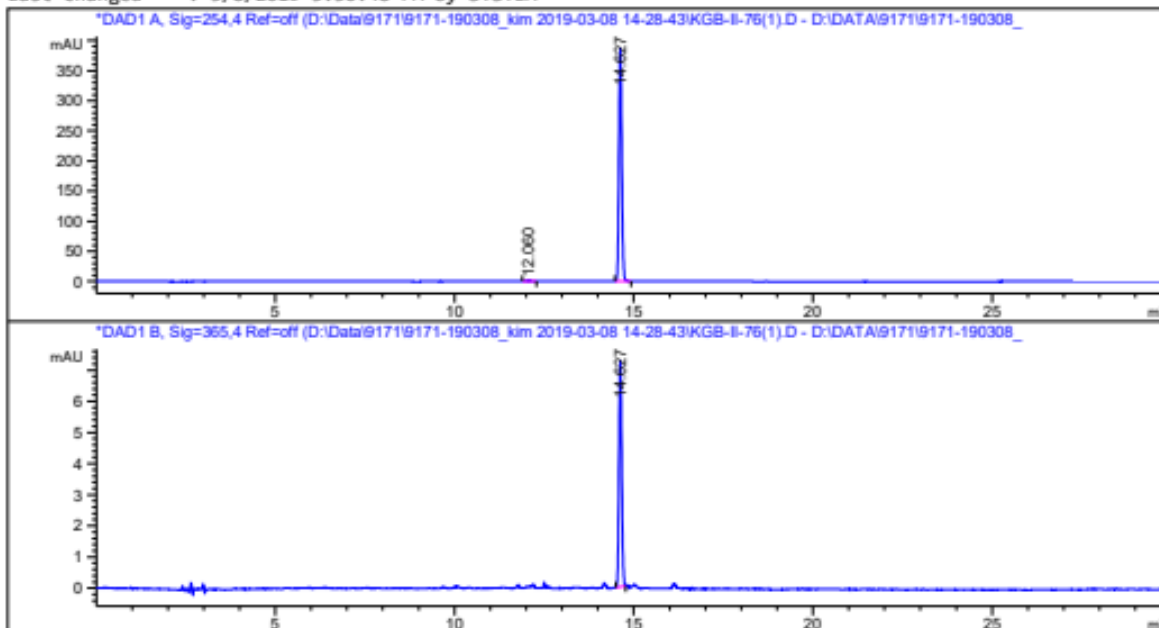
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1	2.748	BB	0.0527	10.08669	3.04008	0.2170
2	2.937	BB	0.1353	15.64708	1.53052	0.3366
3	13.772	BB	0.0918	22.83916	3.81585	0.4913
4	14.661	BB	0.0903	4589.68164	761.54034	98.7254
5	15.311	BB	0.0935	10.68061	1.74166	0.2297

Totals : 4648.93519 771.66846

HPLC data of the compound **5g**

Data File D:\Data\9171\9171-190308_kim 2019-03-08 14-28-43\KGB-II-76(1).D
Sample Name: KGB-II-76(1)

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Injection Date  : 3/8/2019 6:36:49 PM         Inj       :    1
                                                Inj Volume: 10.000 µl
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Last changed    : 3/8/2019 3:14:53 PM by SYSTEM
Analysis Method : D:\Data\9171\9171-190308_kim 2019-03-08 14-28-43\190308_kim.M (Sequence
Method)
Last changed    : 3/8/2019 3:30:48 PM by SYSTEM
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```
=====
                          Area Percent Report
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Multiplier     :      1.0000
Dilution       :      1.0000
Use Multiplier & Dilution Factor with ISTDs
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Signal 1: DAD1 A, Sig=254,4 Ref=off
Signal has been modified after loading from rawdata file!

Peak #	RetTime [min]	Type	Width [min]	Area [mAU*s]	Height [mAU]	Area %
1	12.060	BB	0.0941	10.40172	1.63749	0.5011
2	14.627	BB	0.0821	2065.25195	387.94006	99.4989

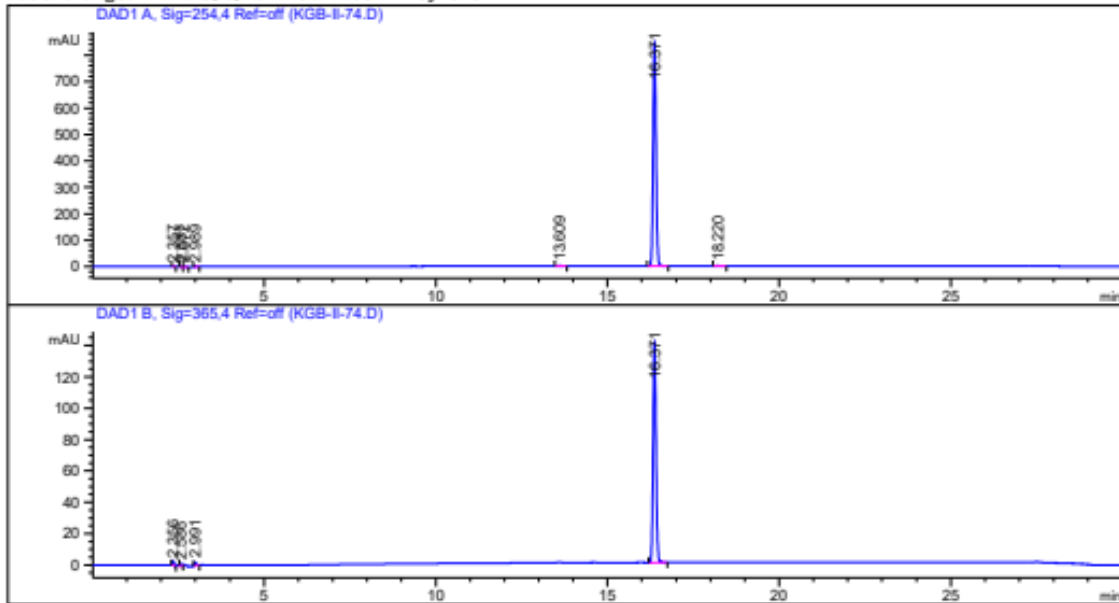
Totals : 2075.65367 389.57755

HPLC data of the compound **5h**

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 Sample Name: KGB-II-74

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Injection Date  : 3/8/2019 5:35:04 PM         Inj       :    1
                                           Inj Volume: 10.000 µl
Sequence File   : D:\Data\9171\9171-190308_kim 2019-03-08 14-28-43\9171-190308_kim.S
Method          : D:\Data\9171\9171-190308_kim 2019-03-08 14-28-43\190308_kim.M (Sequence
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Area Percent Report

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Sorted By      : Signal
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Dilution       : 1.0000
Use Multiplier & Dilution Factor with ISTDs
  
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Signal 1: DAD1 A, Sig=254,4 Ref=off

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1	2.357	BB	0.0506	9.57488	2.89289	0.1960
2	2.587	VV	0.0674	5.21559	1.13854	0.1067
3	2.672	VB	0.0747	5.29344	1.05193	0.1083
4	2.989	BB	0.0503	10.31255	3.13893	0.2111
5	13.609	BB	0.0929	10.03435	1.56261	0.2054
6	16.371	BB	0.0867	4836.85352	845.49213	98.9963
7	18.220	BB	0.0974	8.60644	1.33185	0.1761

Totals : 4885.89077 856.60887

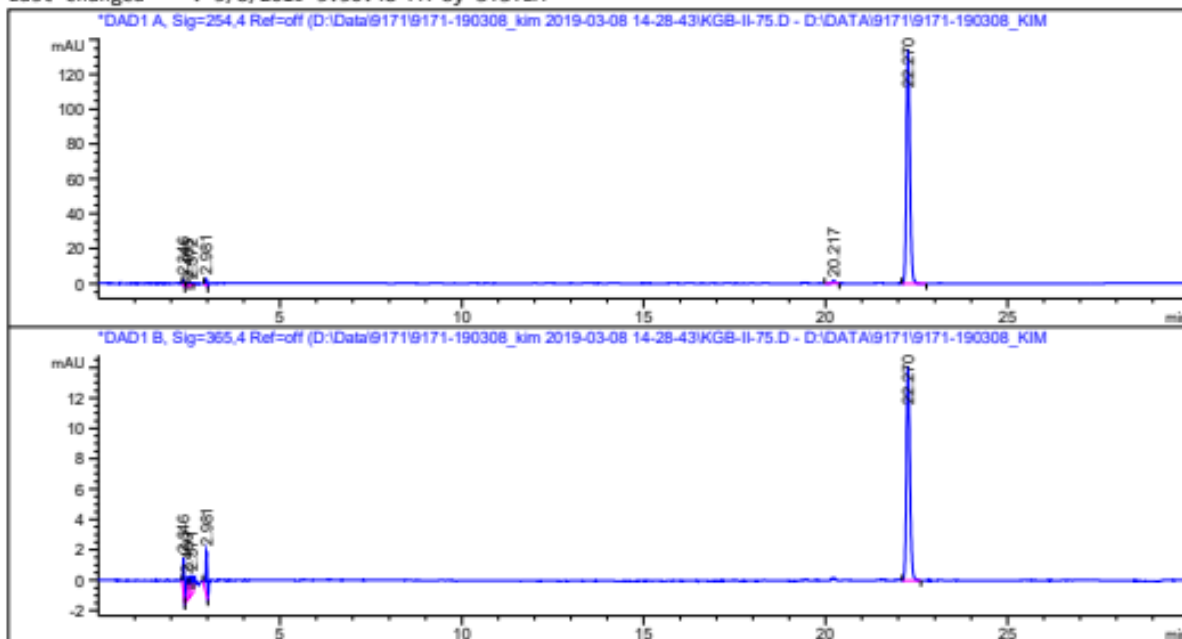
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                                           Inj Volume: 10.000 µl

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Method)
Last changed   : 3/8/2019 3:30:48 PM by SYSTEM
  
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Area Percent Report

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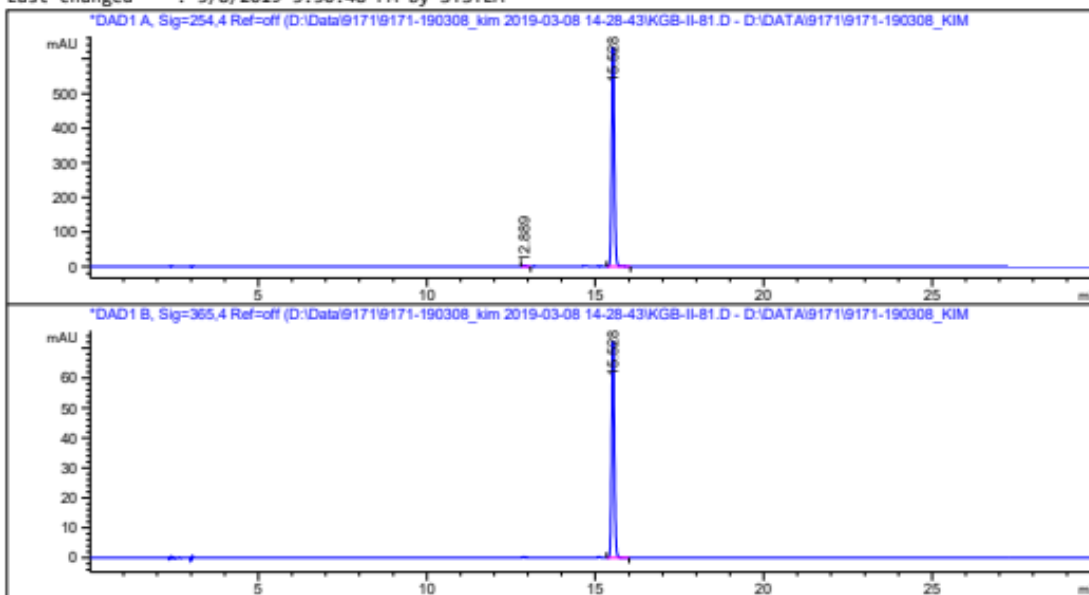
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2	2.463	BV	0.0003	7.73419	1.60179	0.8173
3	2.572	VV	0.0785	7.98579	1.35988	0.8439
4	2.981	BB	0.0444	10.16790	3.66342	1.0745
5	20.217	BB	0.0996	8.04234	1.24146	0.8499
6	22.270	BB	0.1046	906.08051	134.56239	95.7504

HPLC data of the compound *Sj*

Data File D:\Data\9171\9171-190308_kim 2019-03-08 14-28-43\KGB-II-81.D
Sample Name: KGB-II-81

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                                           Inj Volume: 10.000 µl
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Analysis Method: D:\Data\9171\9171-190308_kim 2019-03-08 14-28-43\190308_kim.M (Sequence
Method)
Last changed   : 3/8/2019 3:30:48 PM by SYSTEM
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Area Percent Report
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Multiplier     : 1.0000
Dilution       : 1.0000
Use Multiplier & Dilution Factor with ISTDs
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Signal 1: DAD1 A, Sig=254,4 Ref-off
Signal has been modified after loading from rawdata file!

Peak #	RetTime [min]	Type	Width [min]	Area [mAU*s]	Height [mAU]	Area %
1	12.889	BB	0.0828	15.31868	2.84445	0.4422
2	15.528	BB	0.0833	3449.07275	635.58429	99.5578

Totals : 3464.39143 638.42874

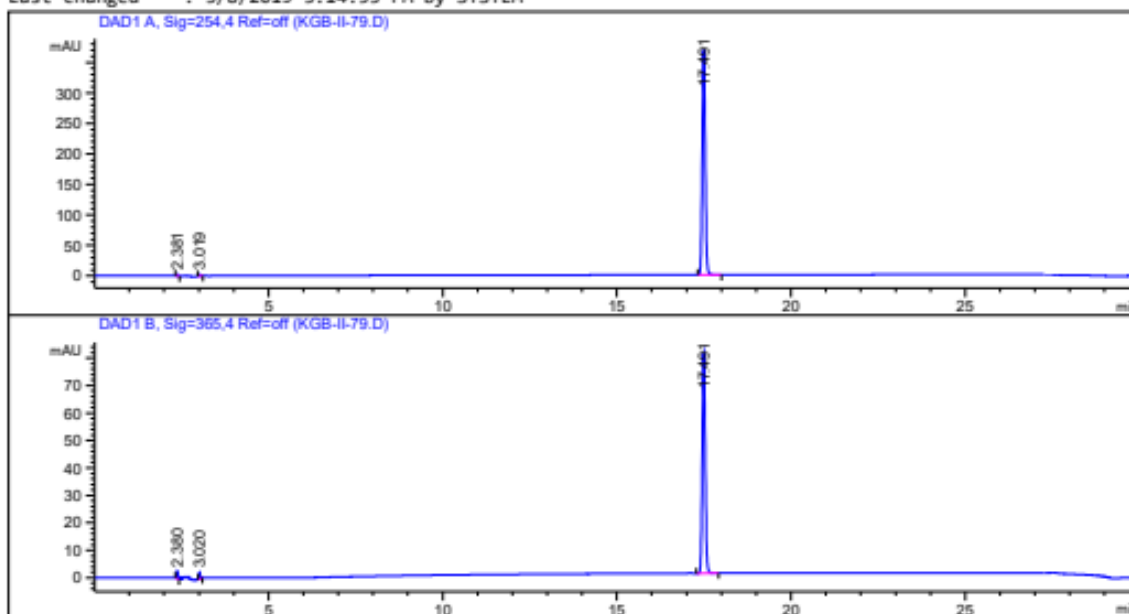
HPLC data of the compound **5k**

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 Sample Name: KGB-II-79

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Injection Date  : 3/8/2019 7:38:34 PM         Inj       :    1
                                           Inj Volume: 10.000 µl

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Method          : D:\Data\9171\9171-190308_kim 2019-03-08 14-28-43\190308_kim.M (Sequence
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 Area Percent Report
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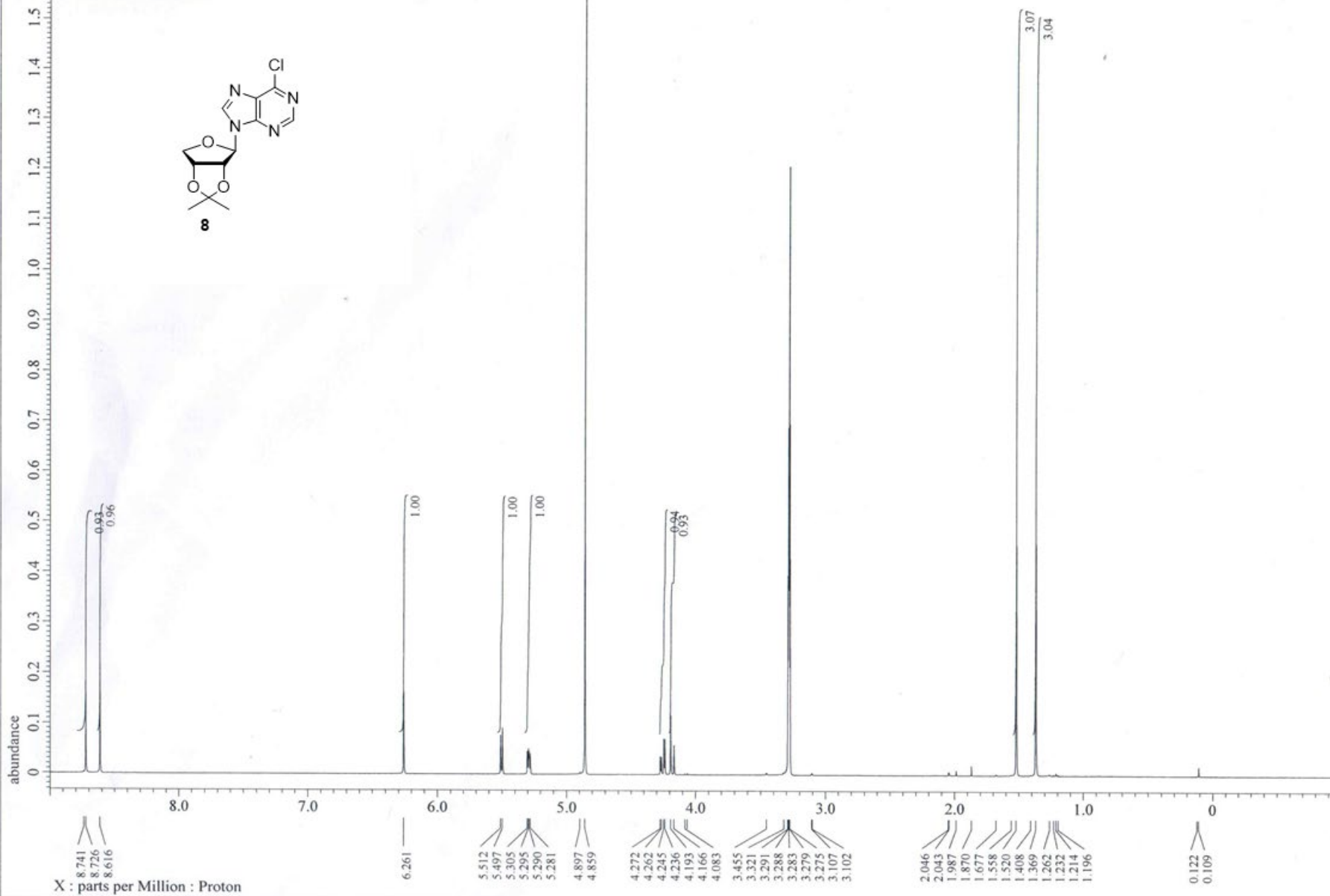
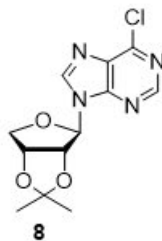
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Dilution       : 1.0000
Use Multiplier & Dilution Factor with ISTDs
  
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Signal 1: DAD1 A, Sig=254,4 Ref=off

Peak #	RetTime [min]	Type	Width [min]	Area [mAU*s]	Height [mAU]	Area %
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2	3.019	BB	0.0474	9.00926	2.97110	0.4118
3	17.491	BB	0.0906	2169.41211	368.91376	99.1711

Totals : 2187.54562 374.96475

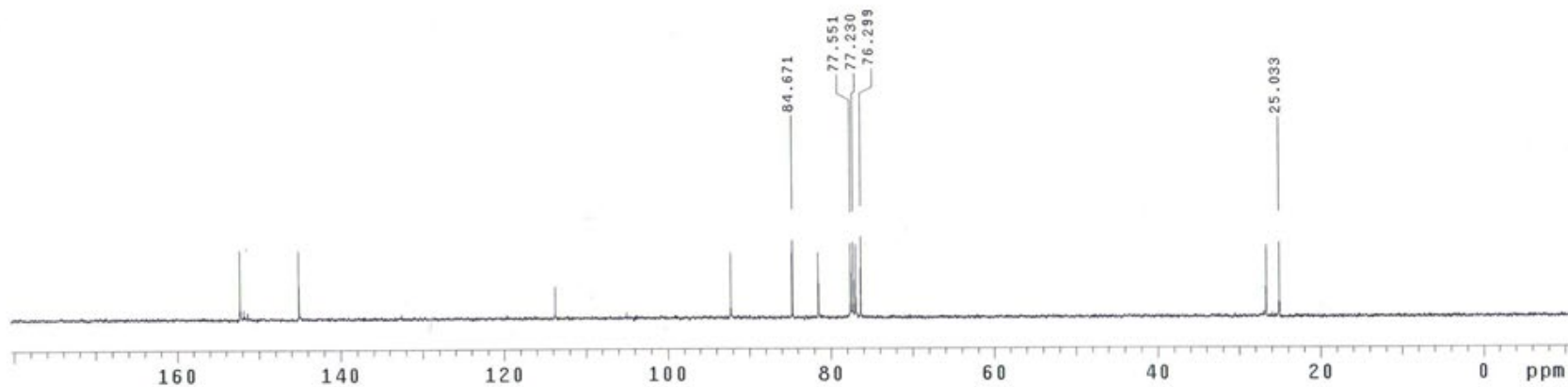
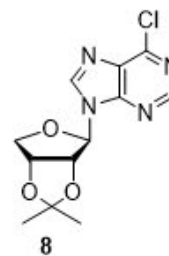
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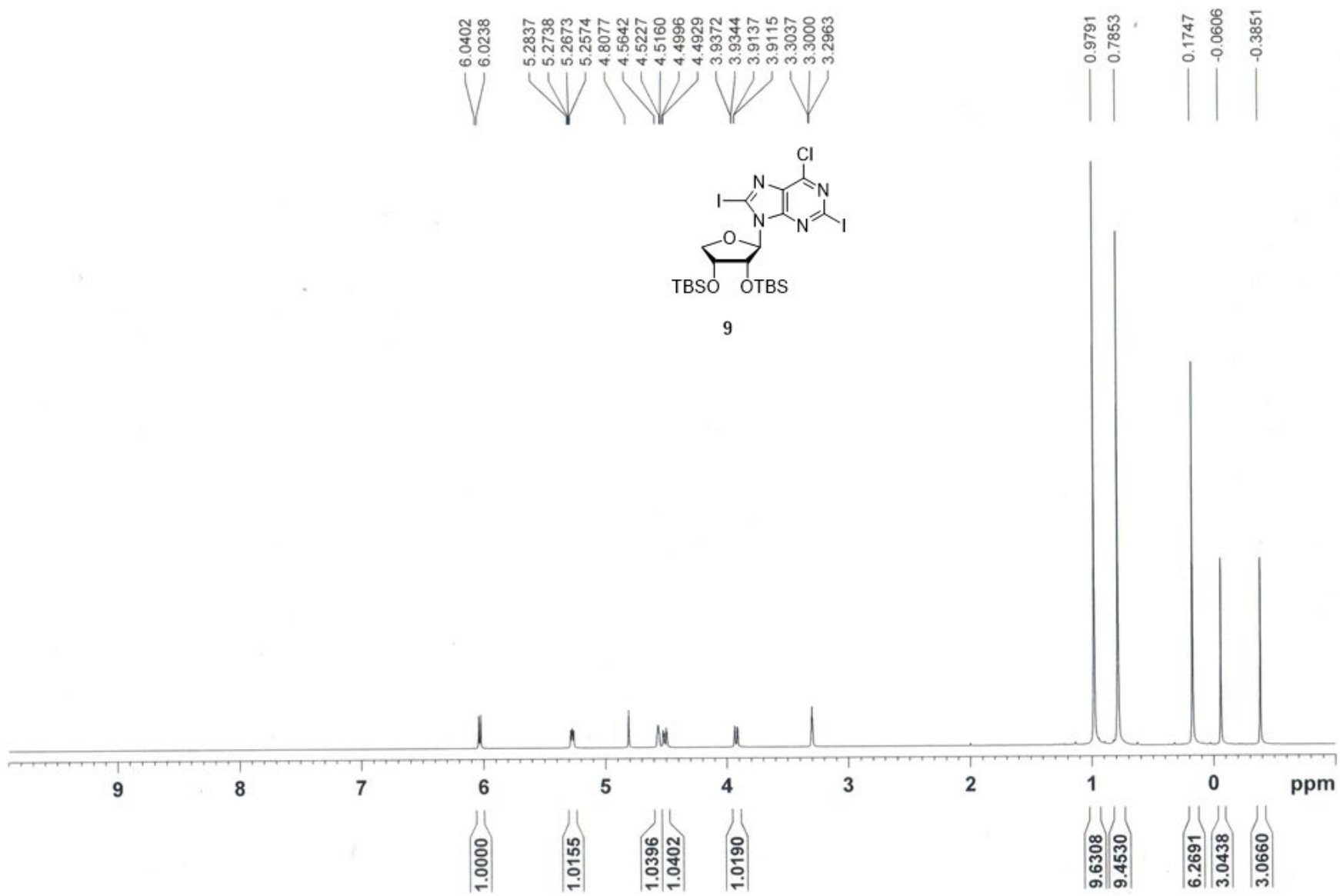


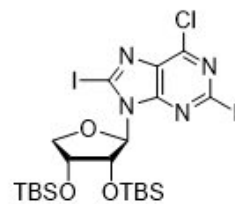
HXY-4-77-1

exp5 std13c

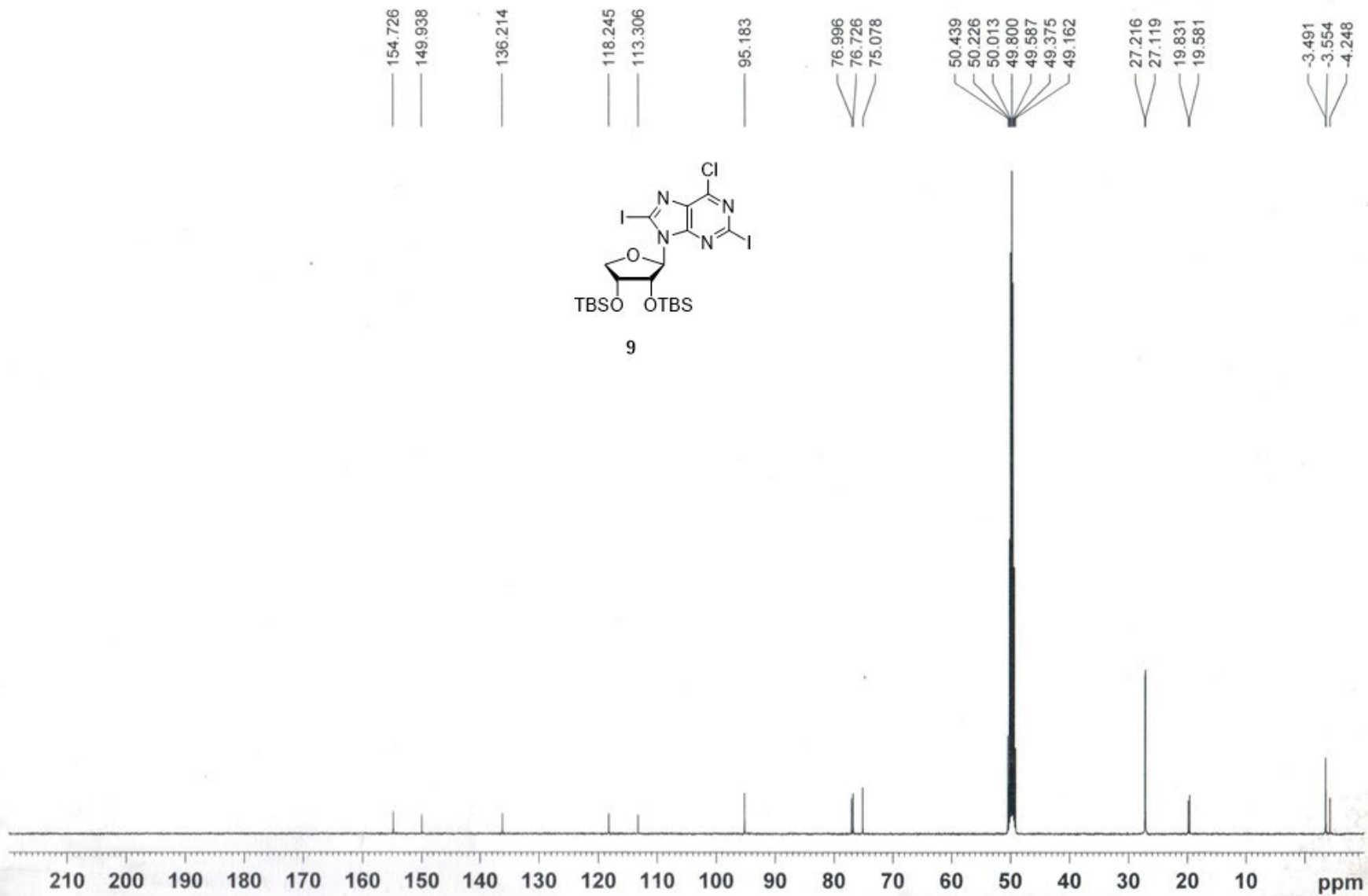
SAMPLE		DEC. & VT	
date	Mar 22 2011	dfrq	400.046
solvent	CDC13	dn	H1
file	exp	dpwr	41
ACQUISITION		dof	1043.9
sfrq	100.601	dm	YYY
tn	C13	dmm	w
at	0.916	daf	10900
np	46076	dseq	
sw	25157.2	dres	1.0
fb	not used	homo	n
bs	4	temp	25.0
tpwr	55	PROCESSING	
pw	10.1	lb	2.00
d1	2.000	wtfile	
tof	1043.3	proc	ft
nt	1500	fn	not used
ct	240	math	f
alock	n		
gain	not used	werr	
FLAGS		wexp	
il	n	wbs	
in	n	wnt	
dp	y		
hs	nn		
DISPLAY			
sp	-1091.8		
wp	19238.7		
vs	264		
sc	0		
wc	250		
hzmm	76.95		
is	41.78		
rfl	9786.4		
rfp	7768.6		
th	12		
ins	1.000		
al	ph		

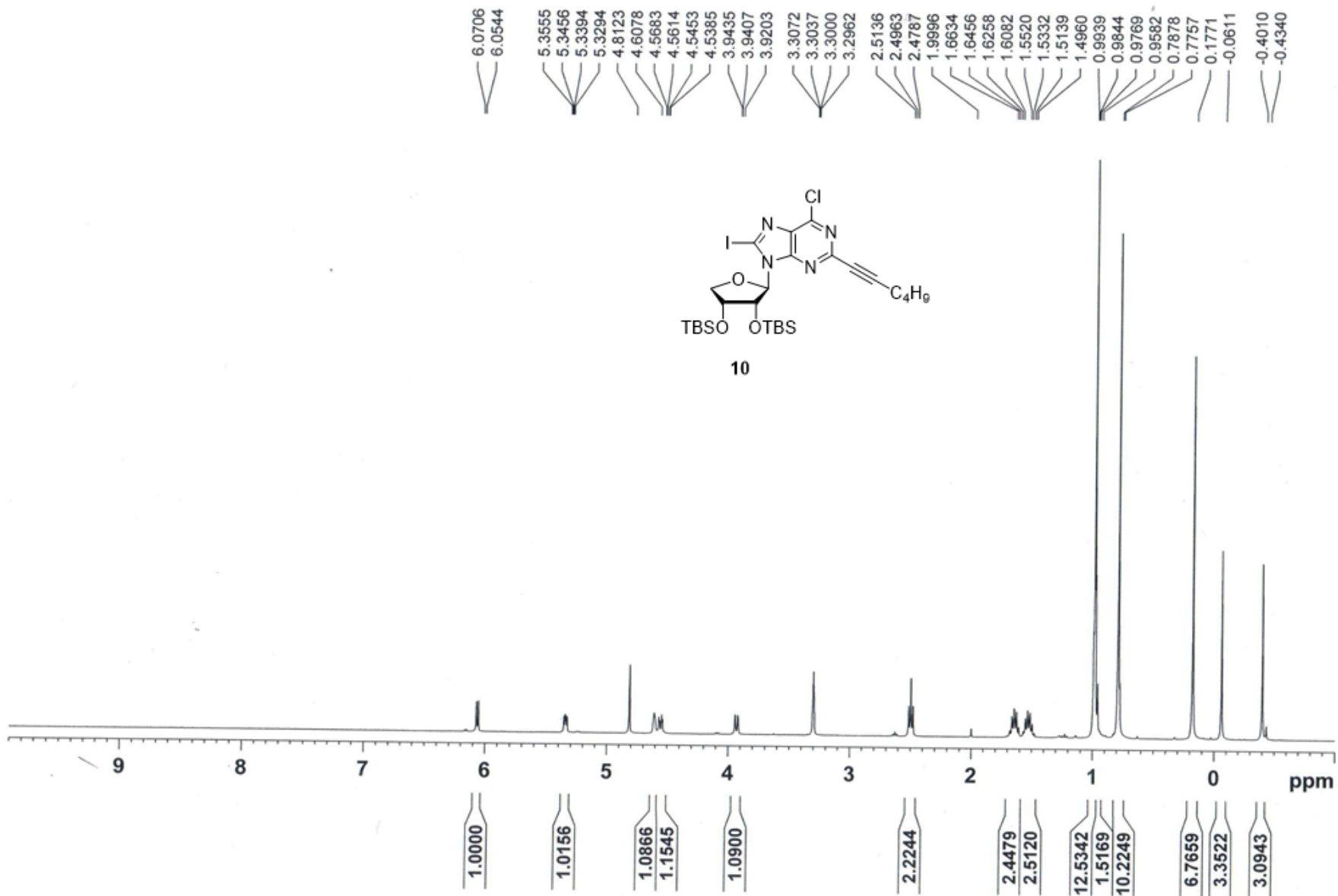


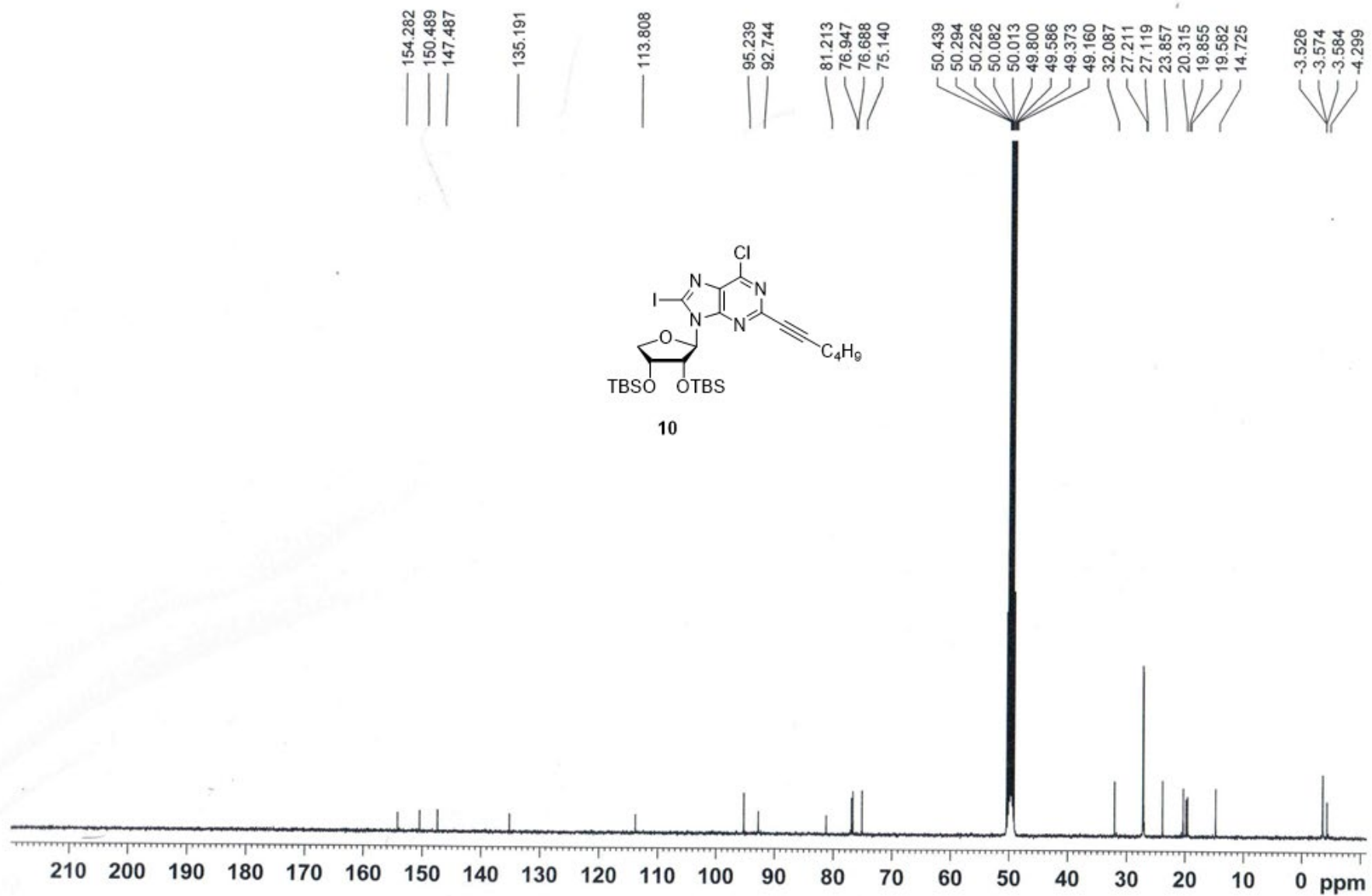


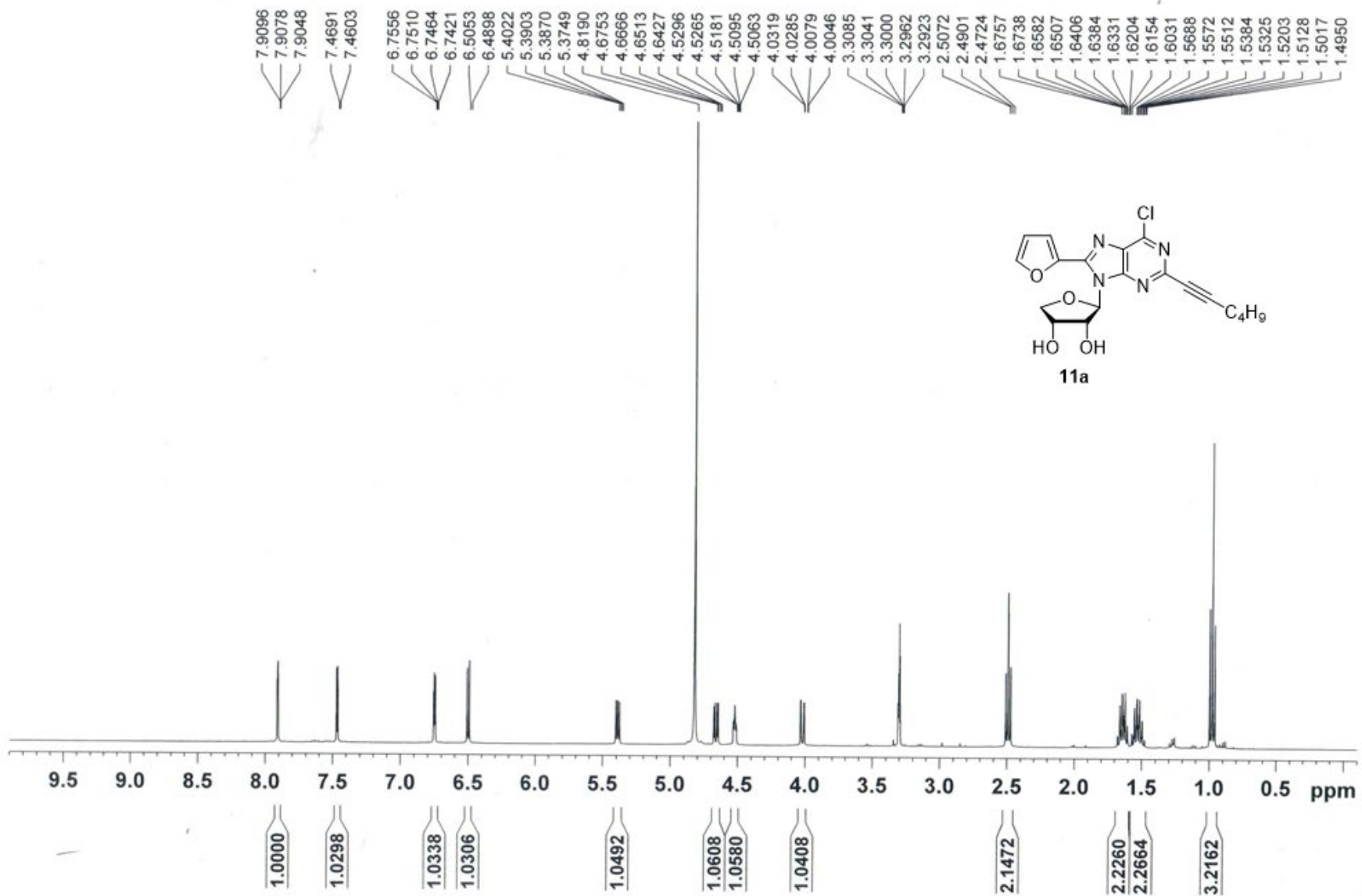


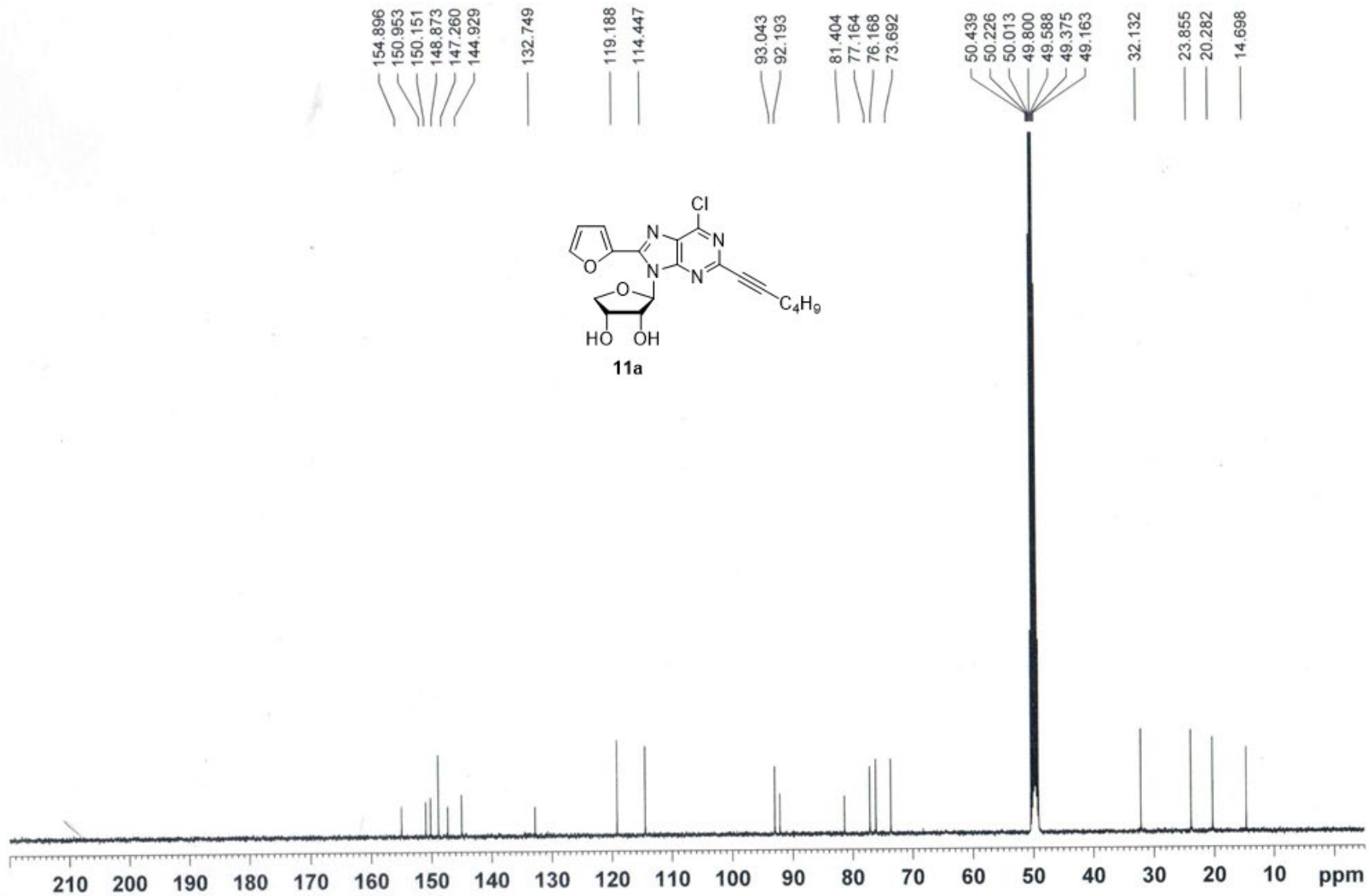
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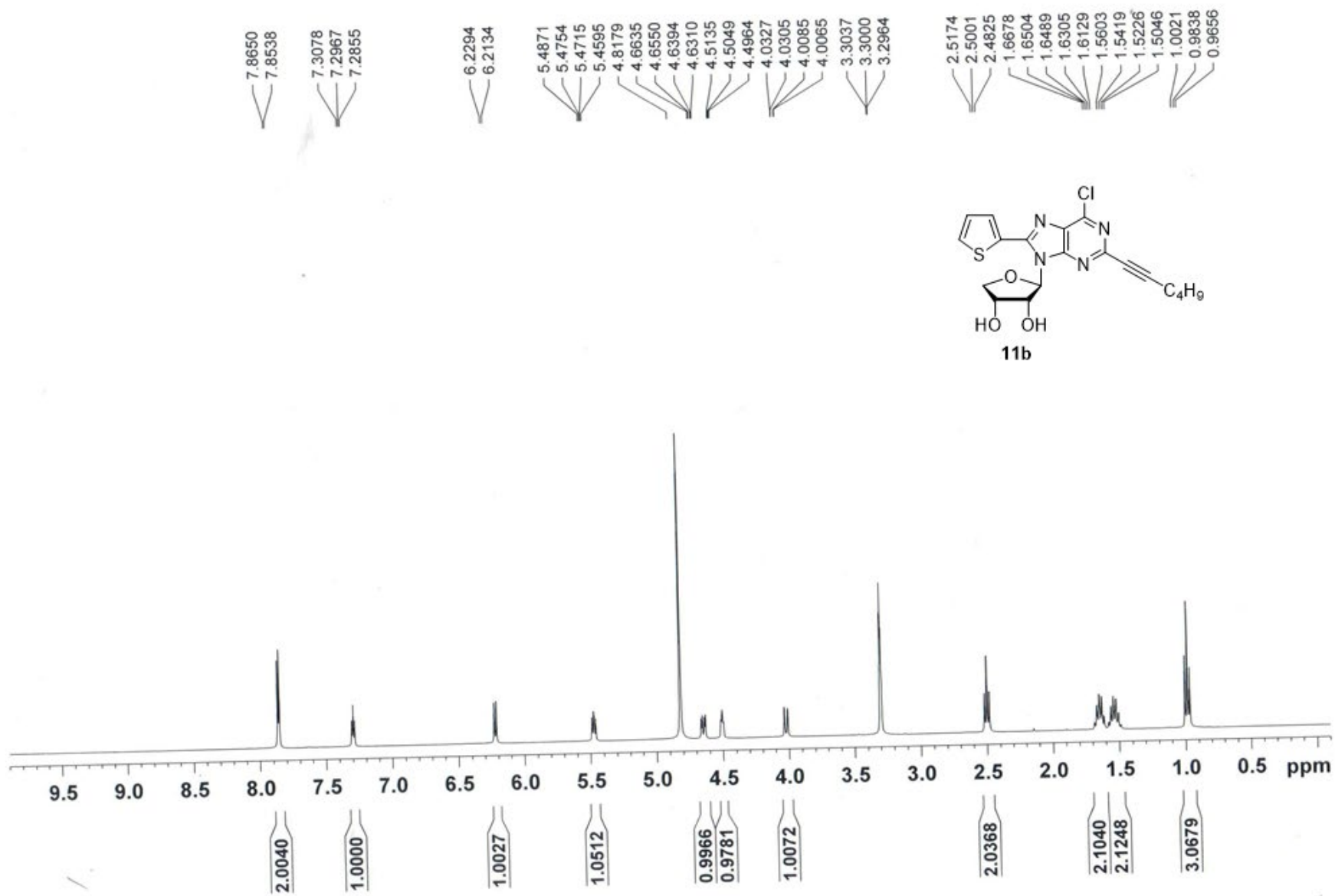


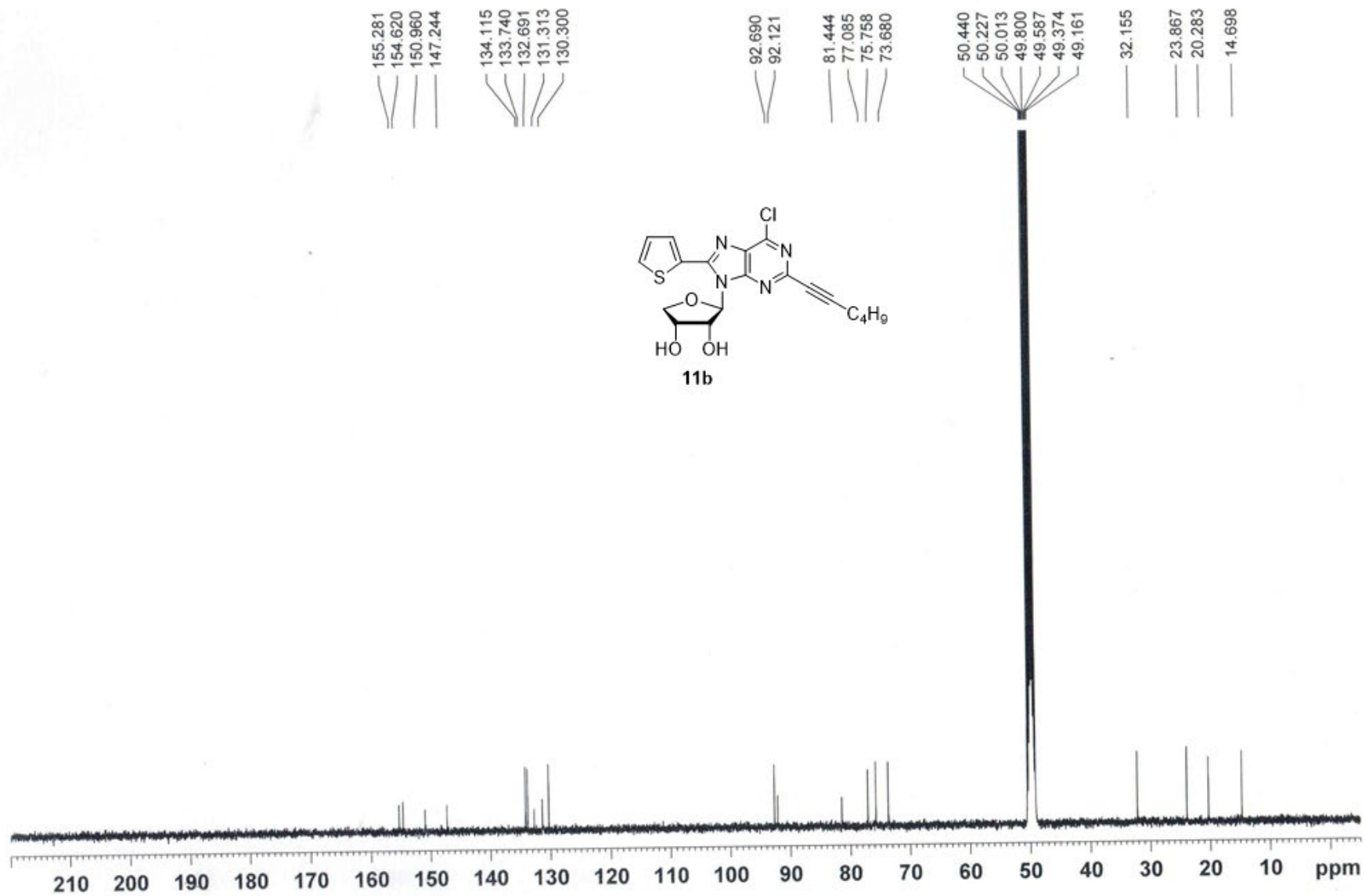


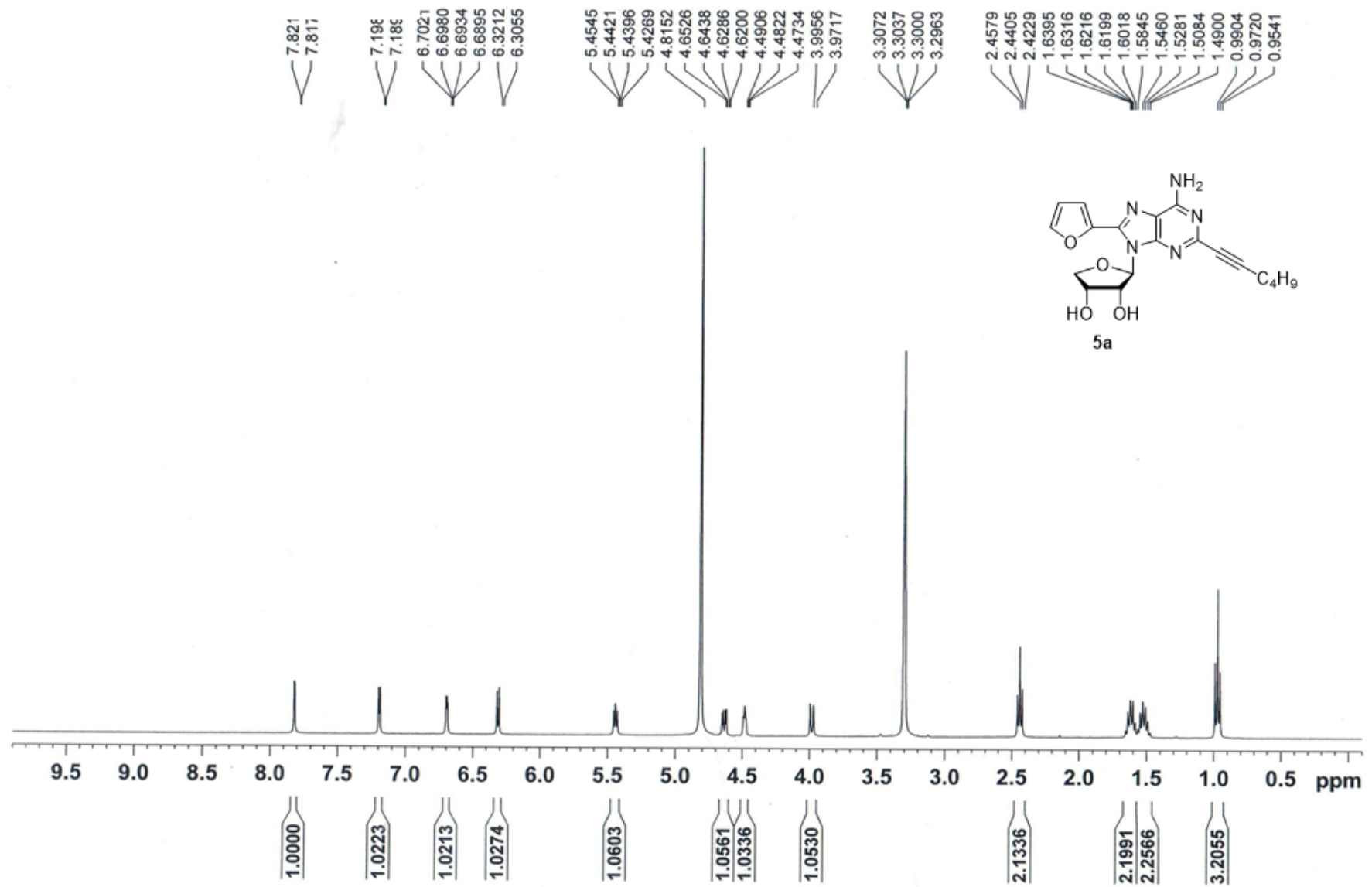


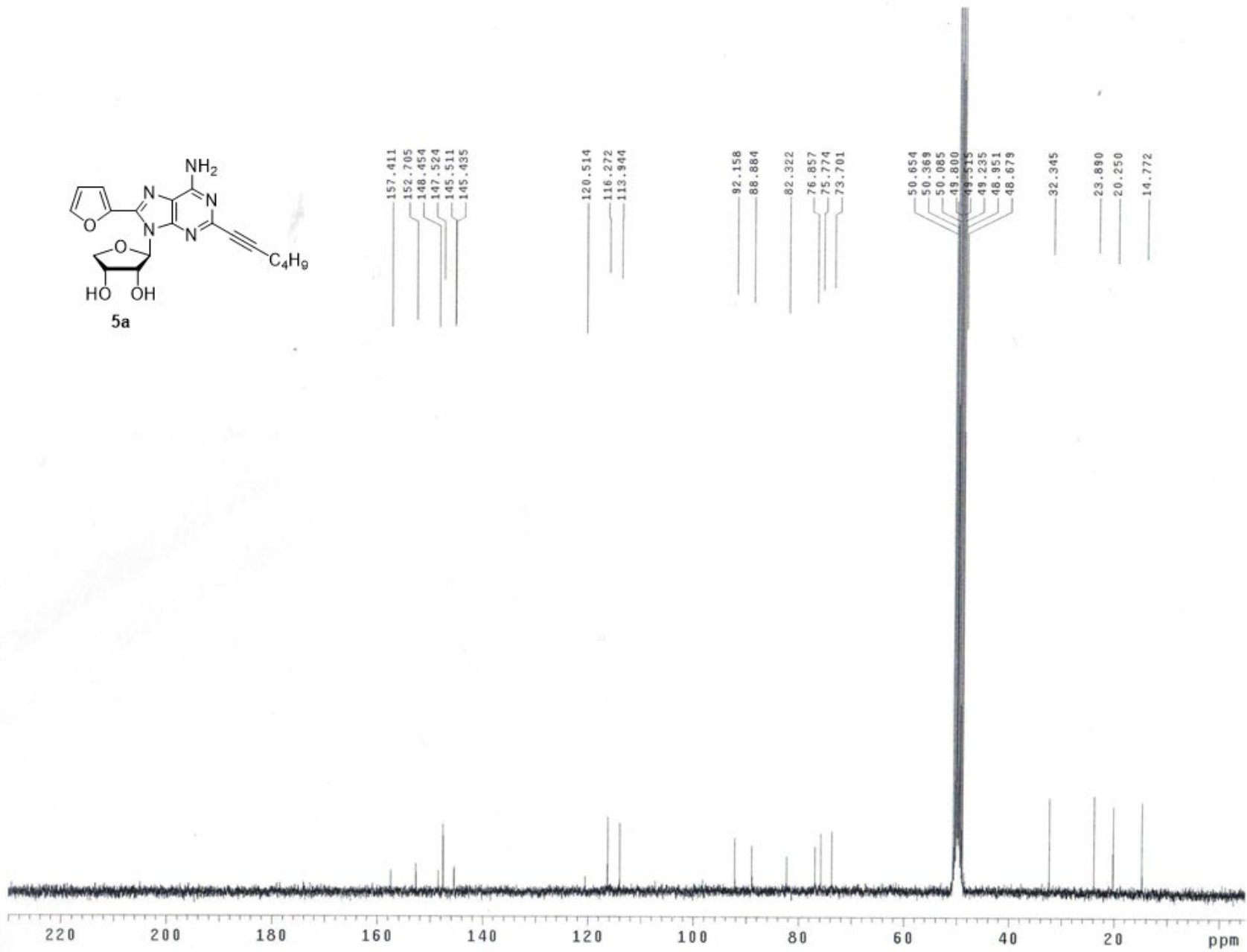


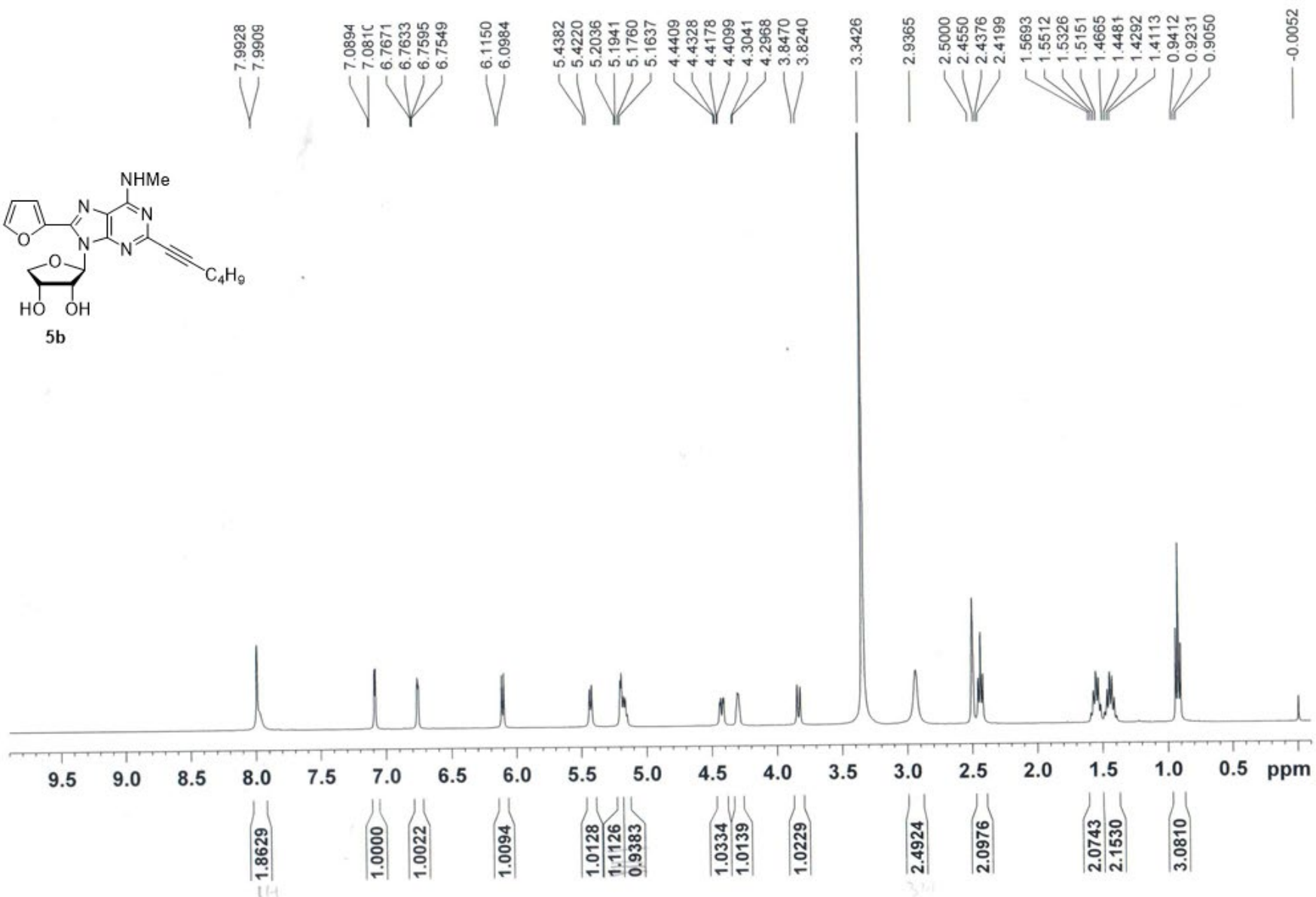
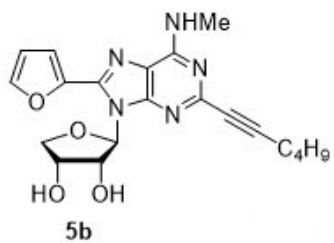


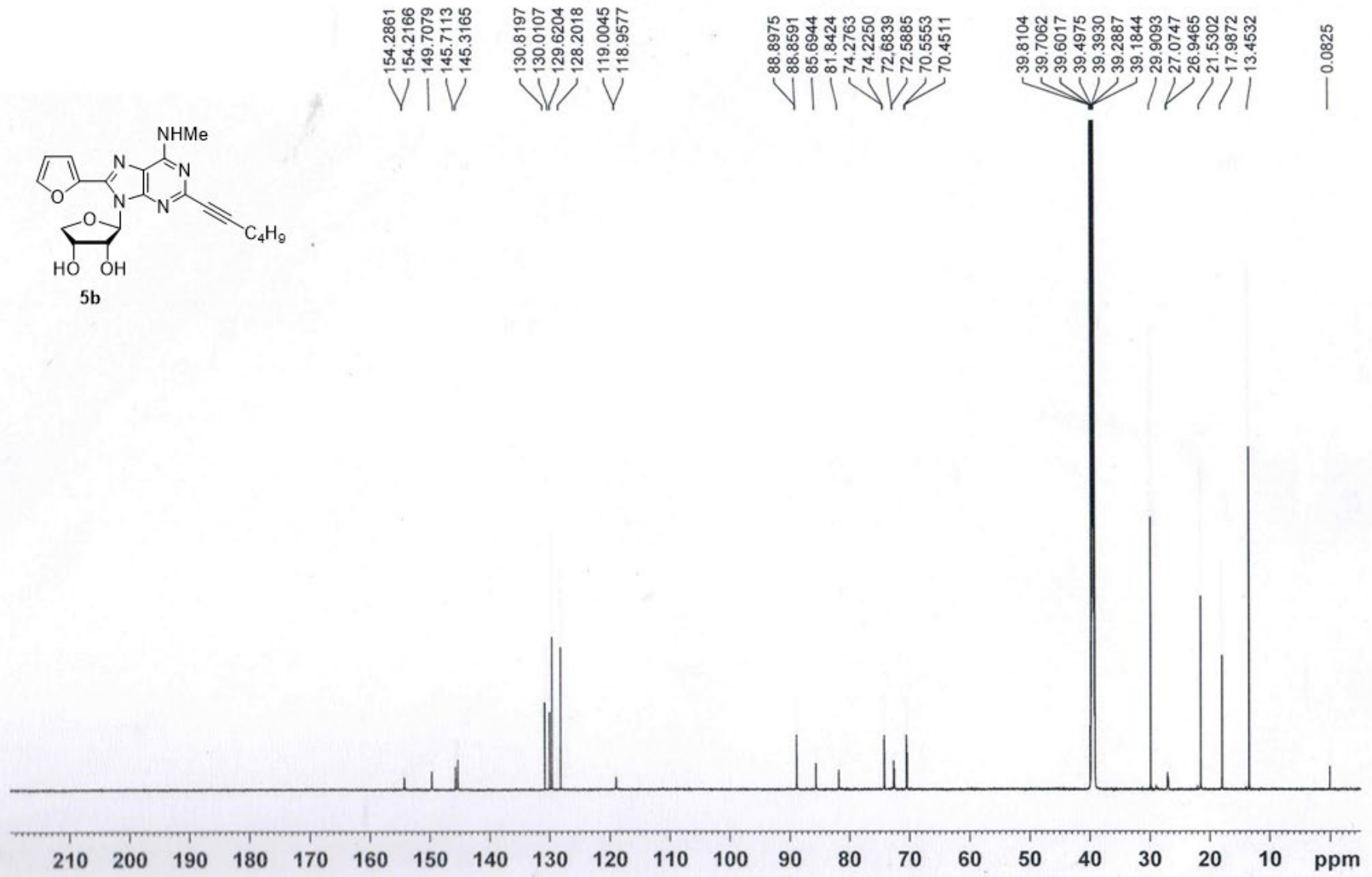
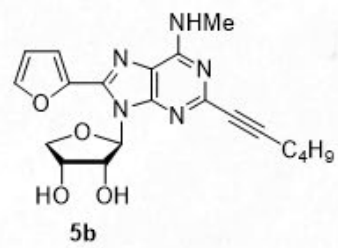


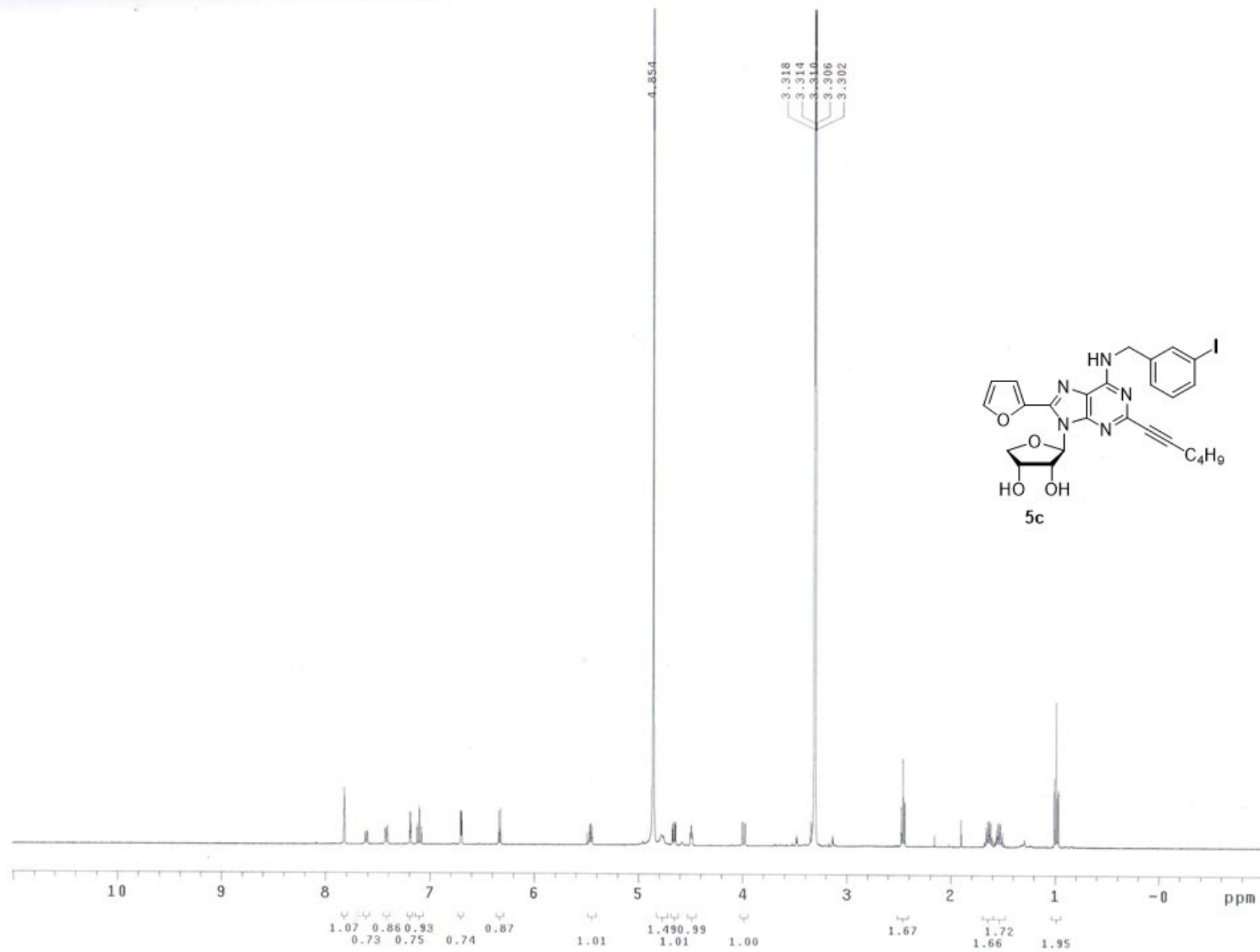


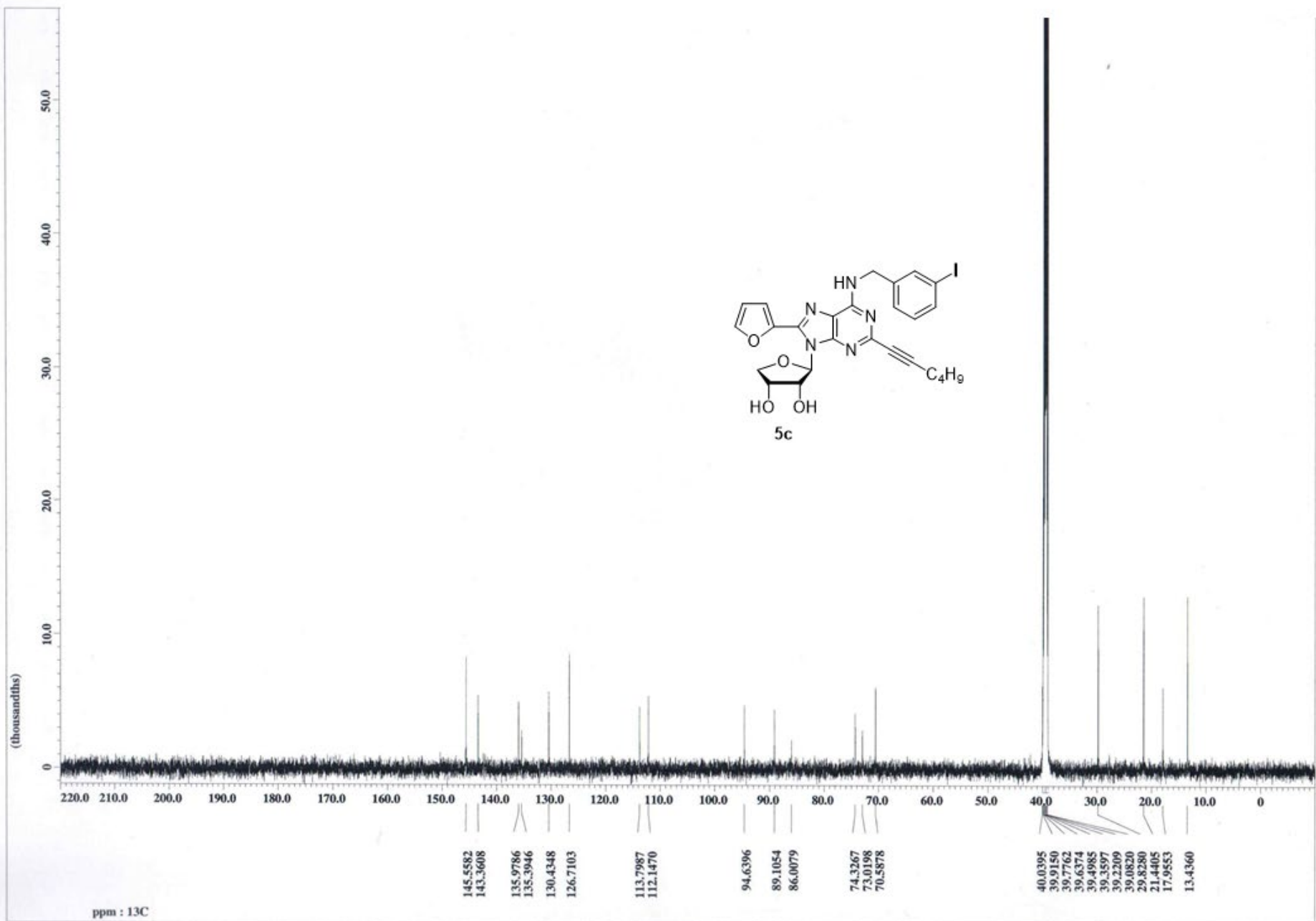


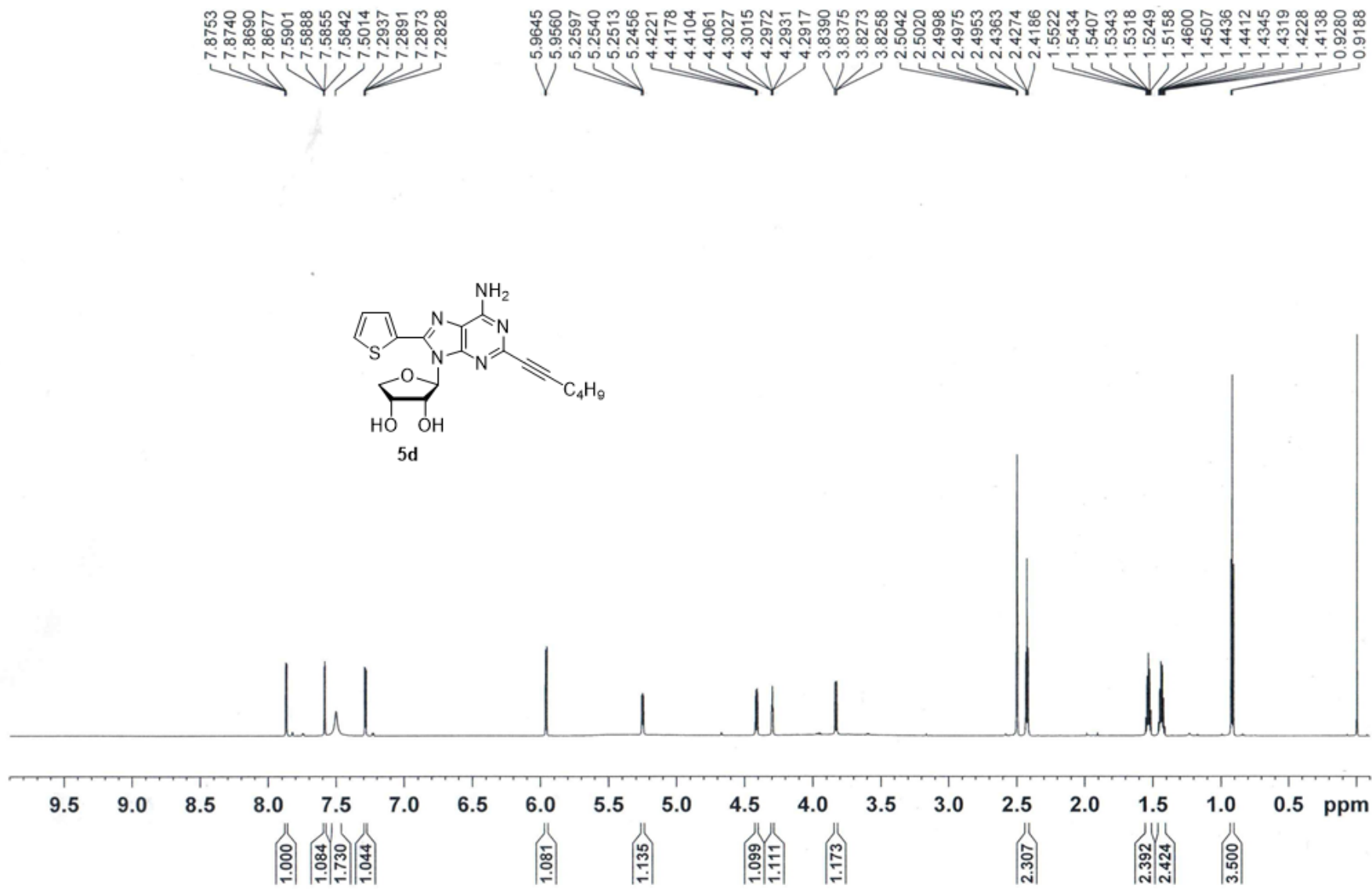


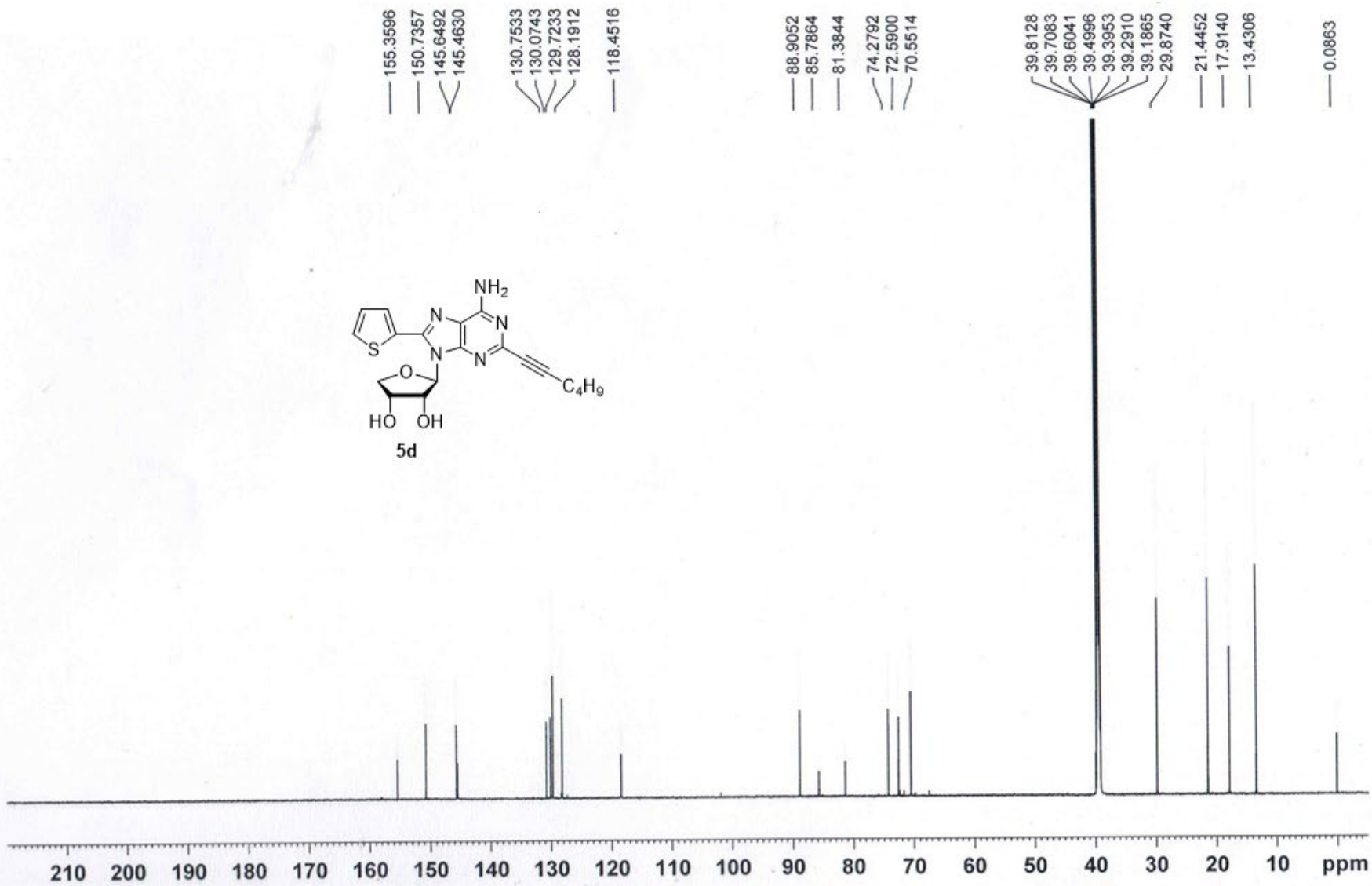


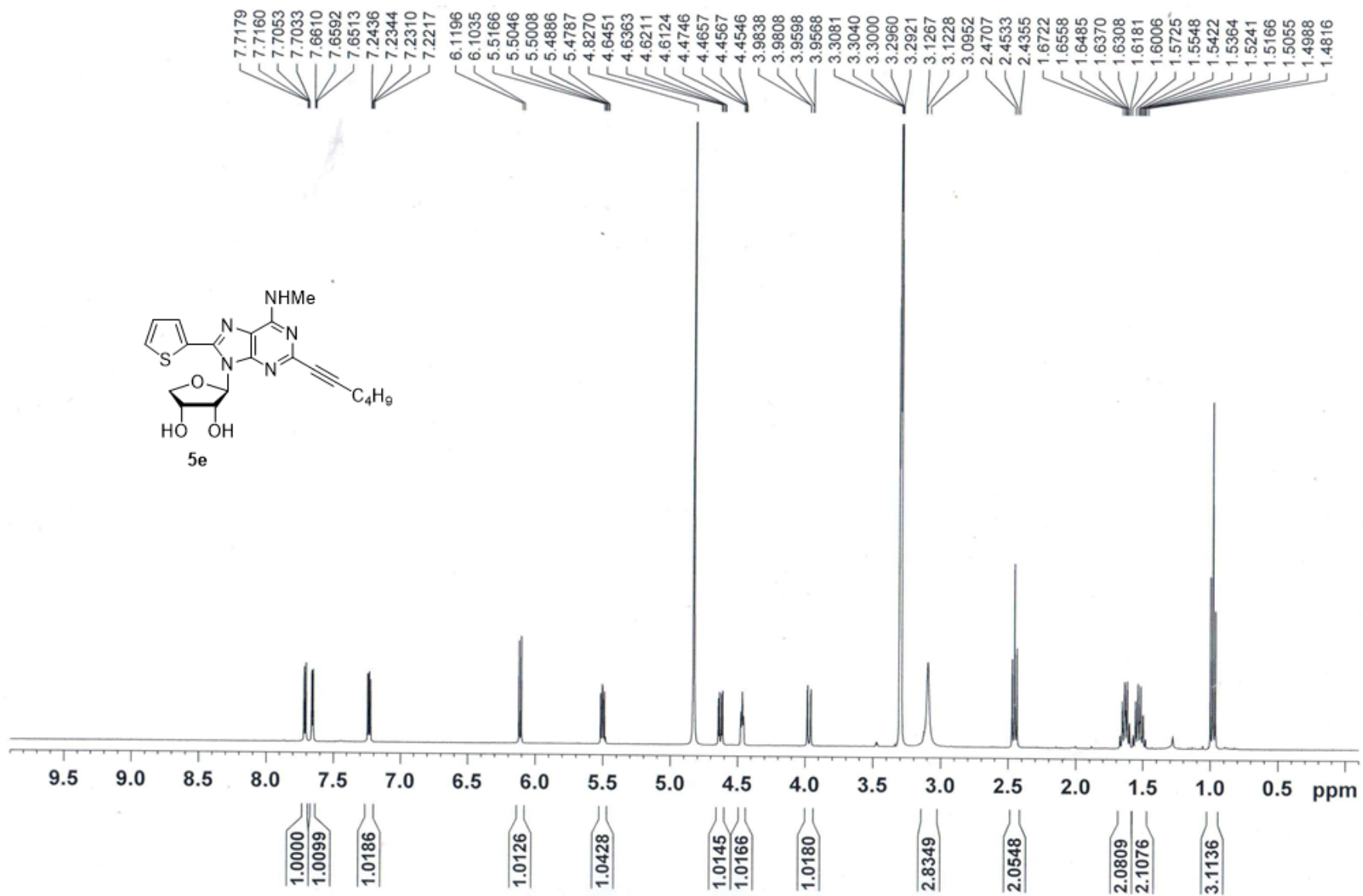


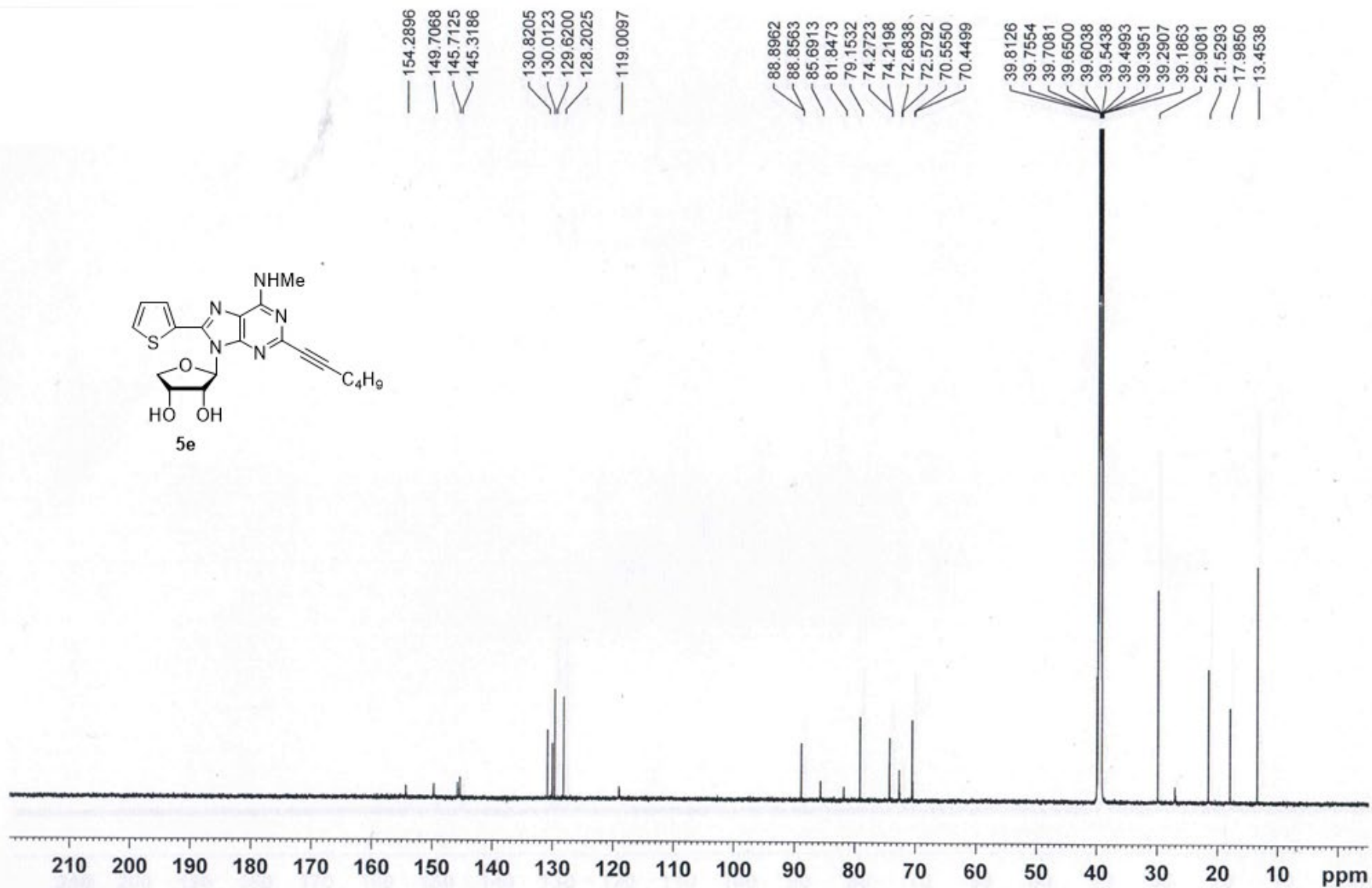
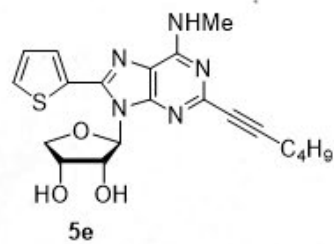


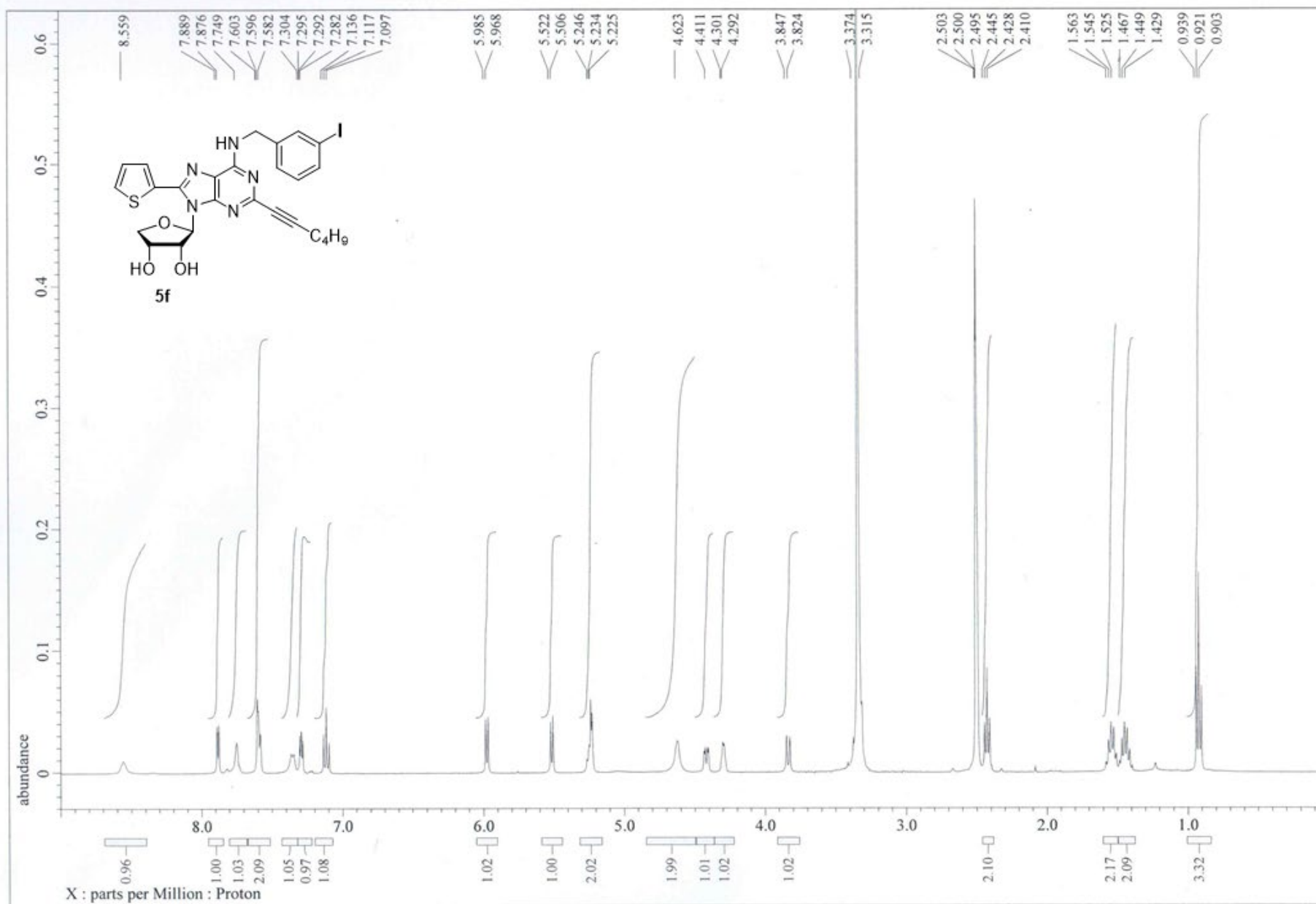


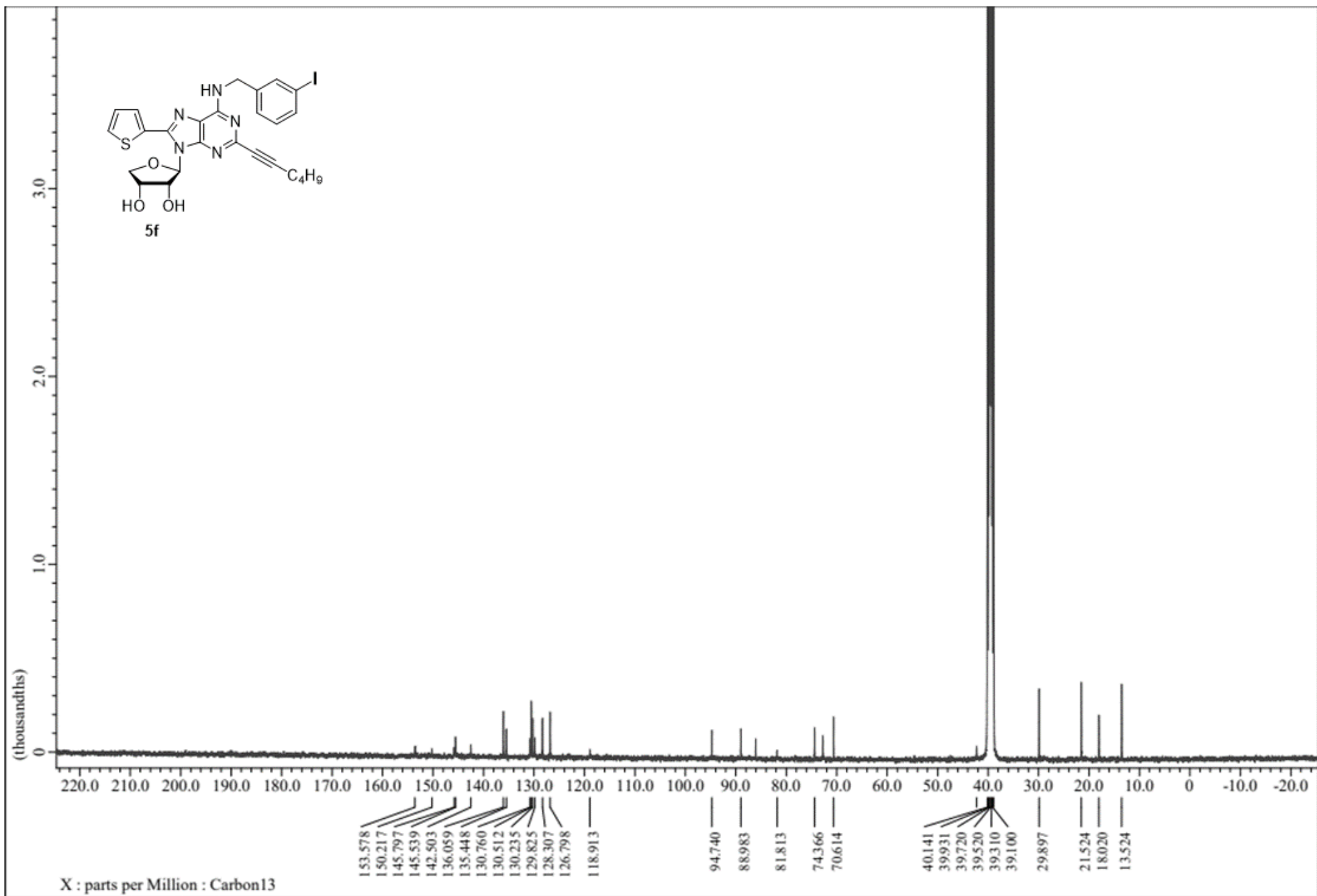


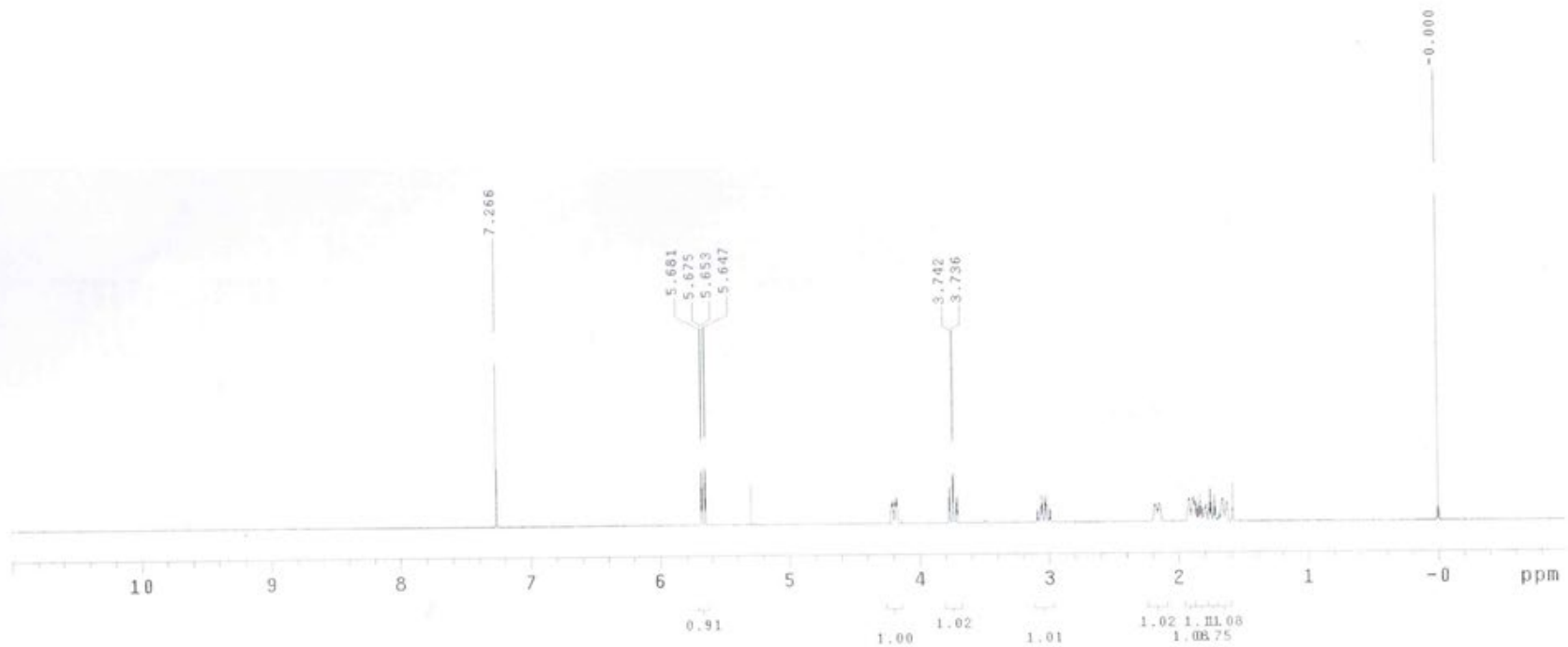
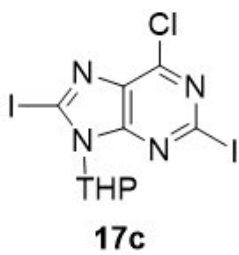


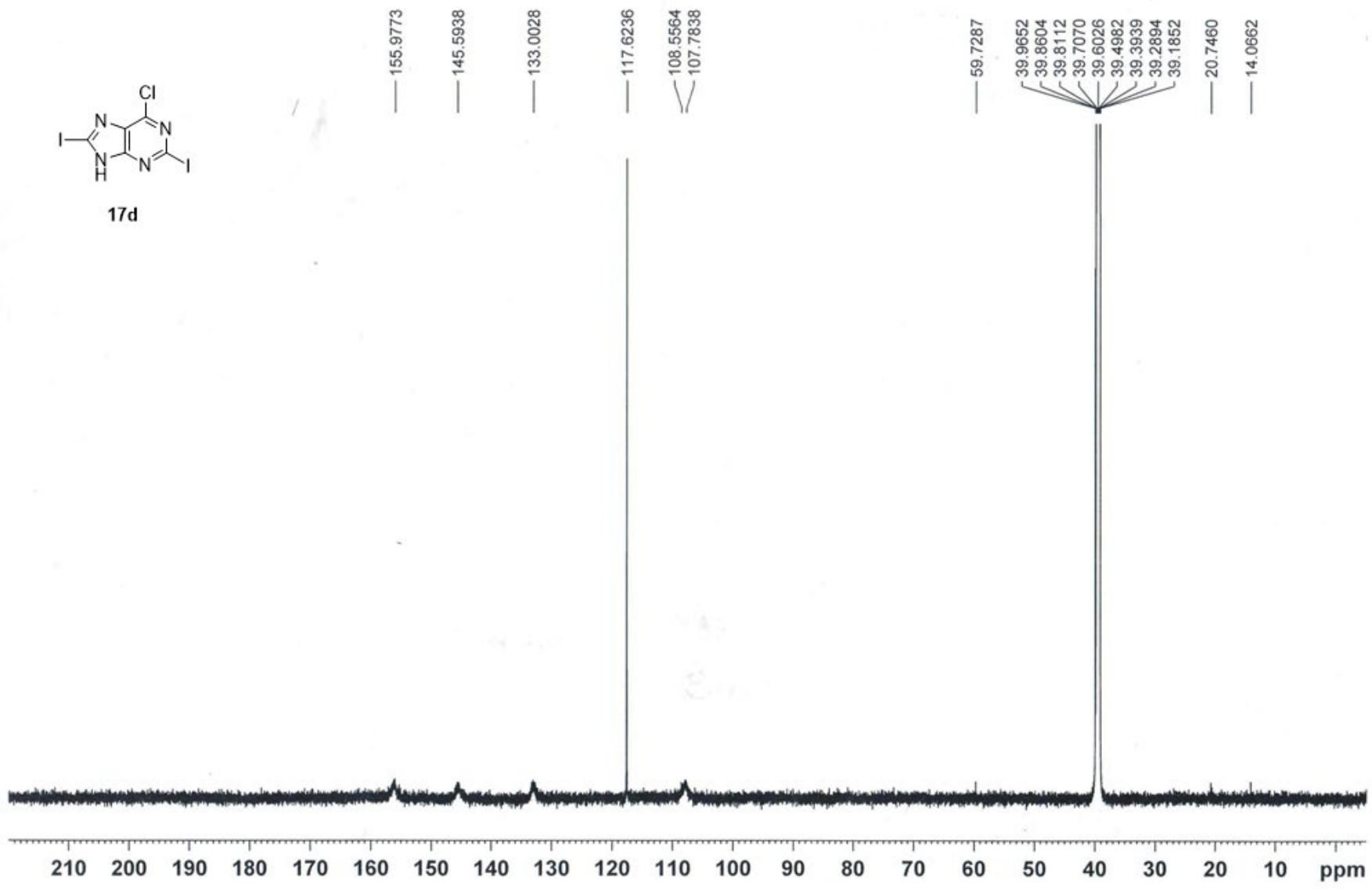




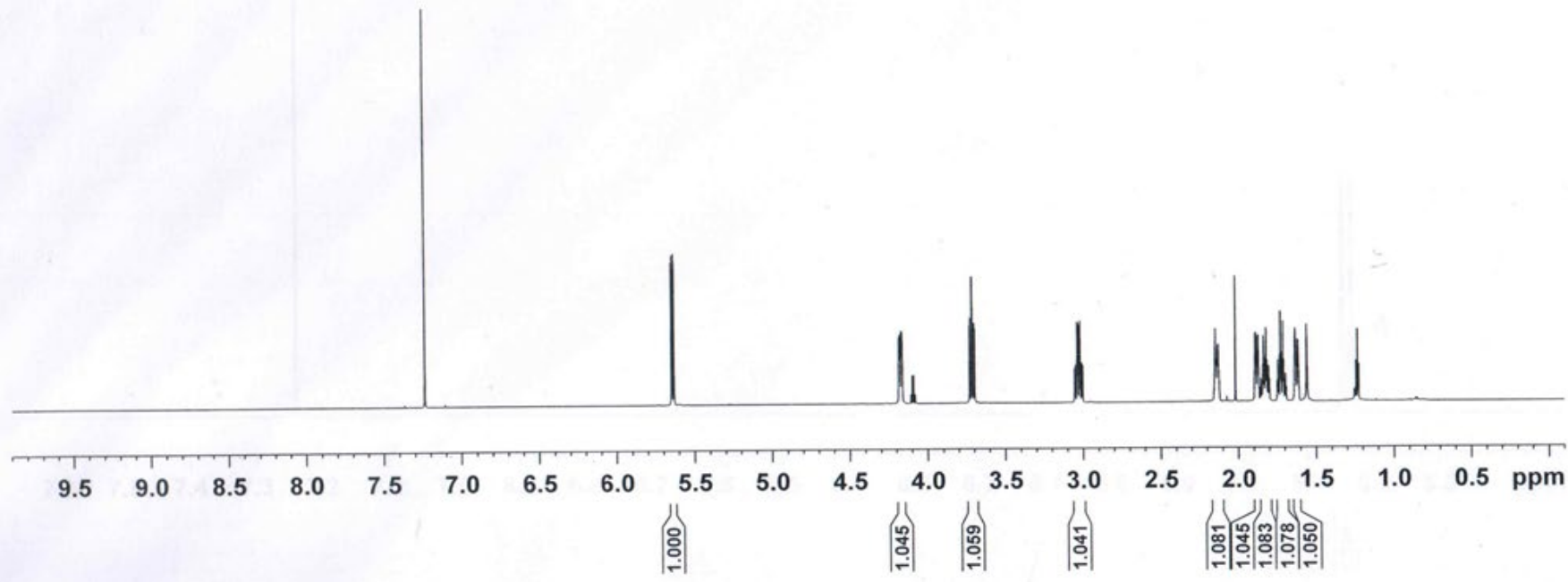
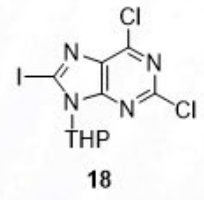


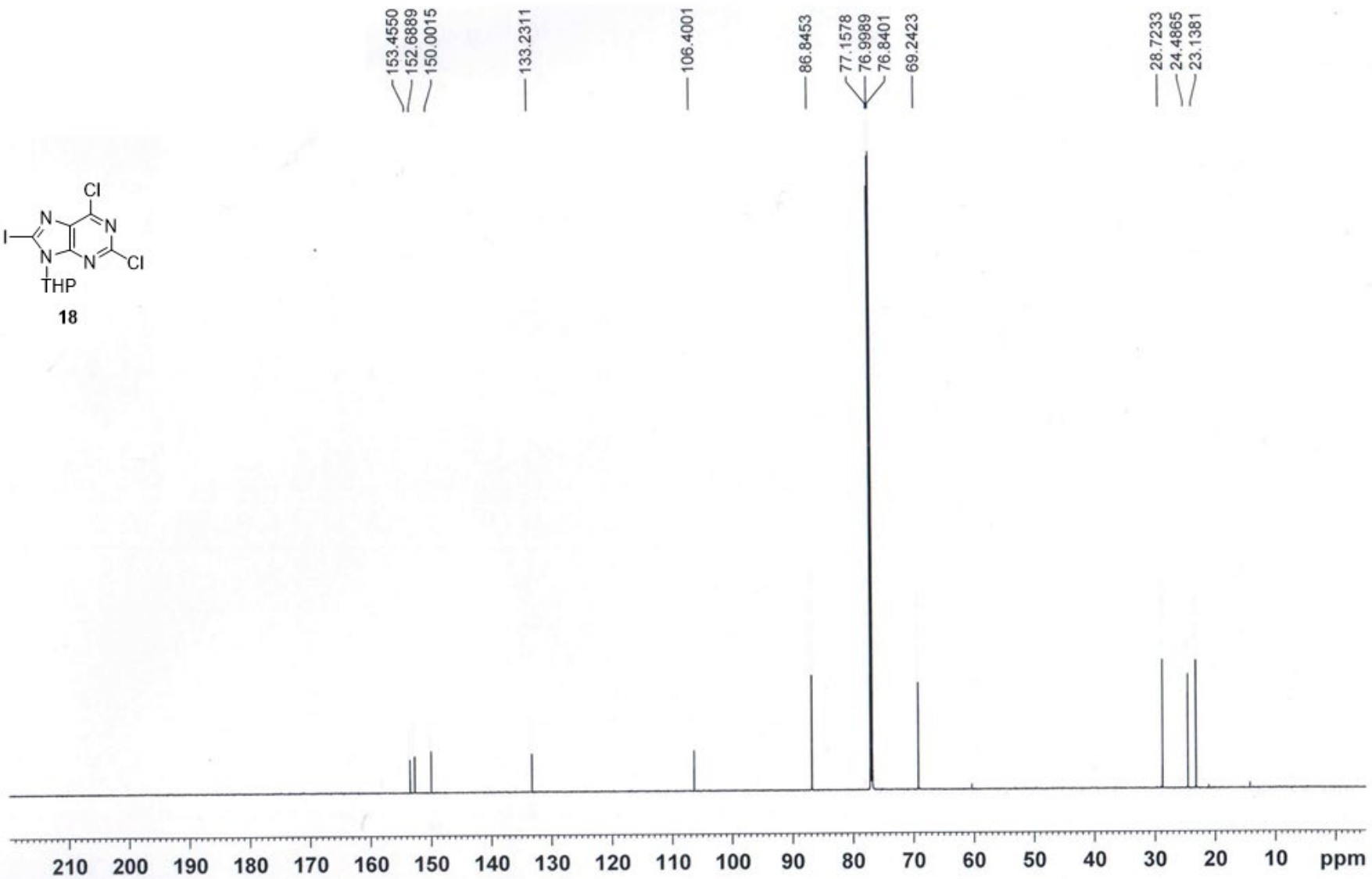
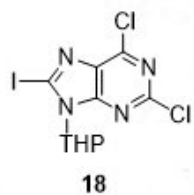


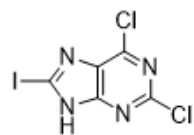




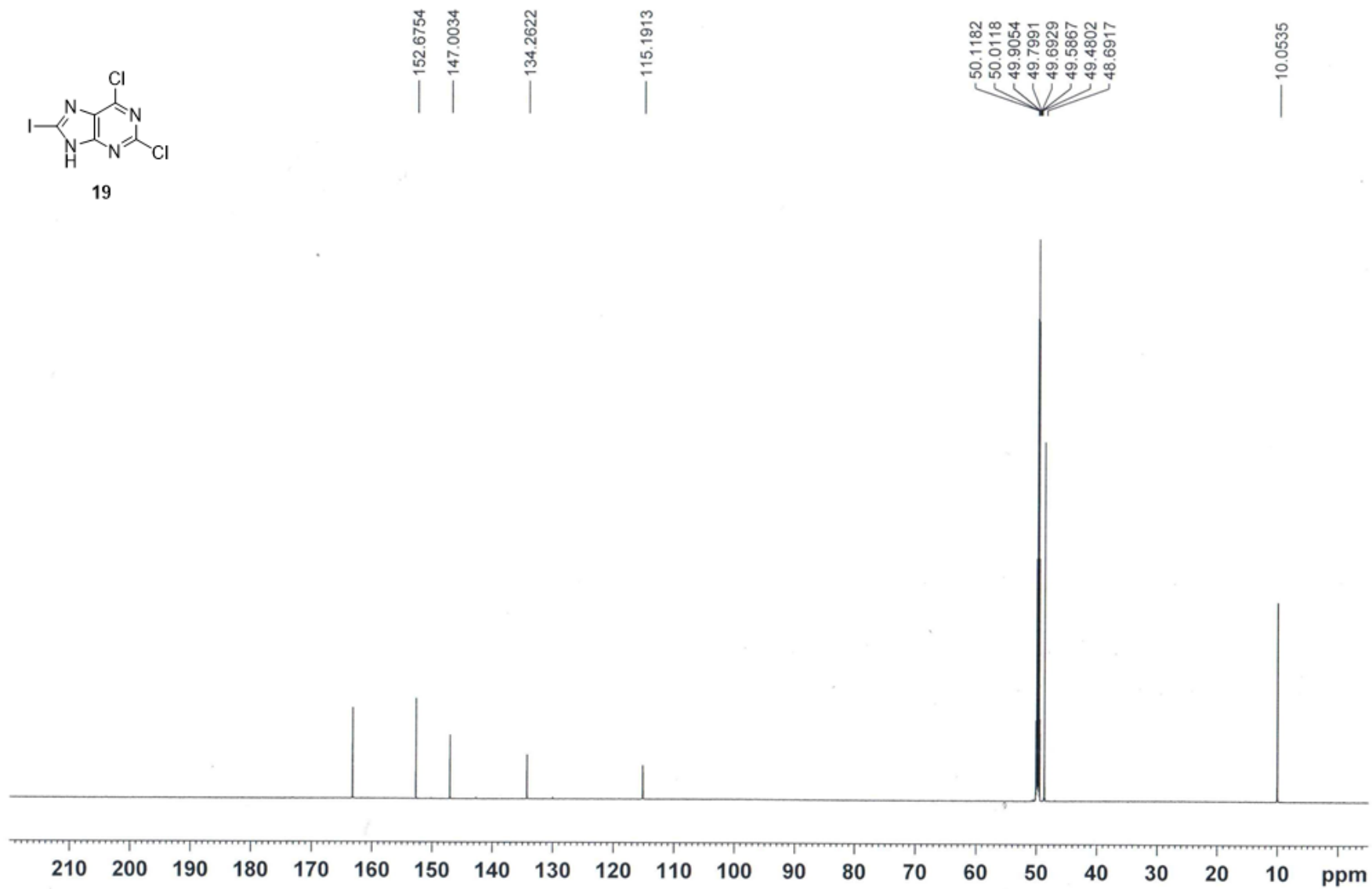
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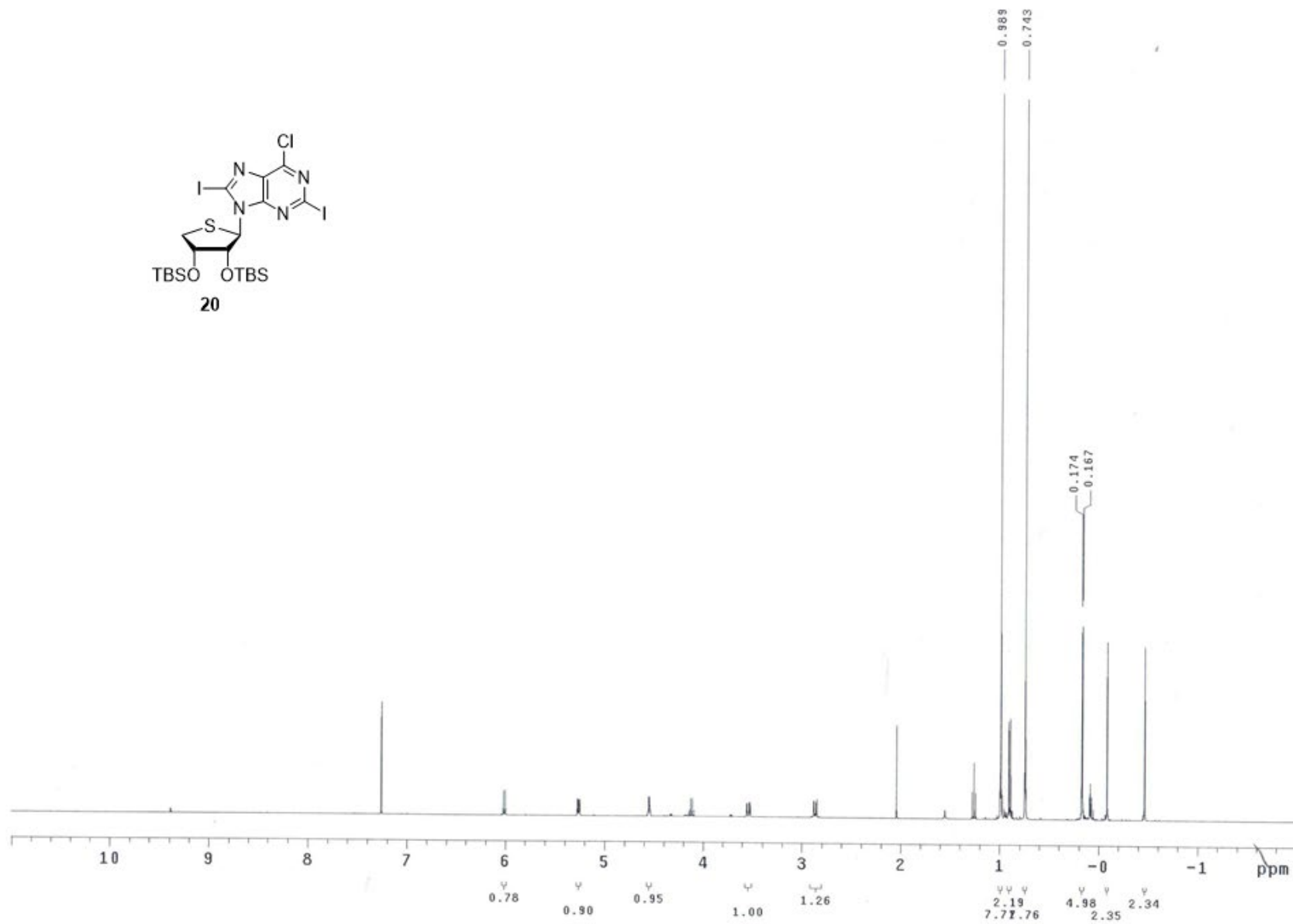
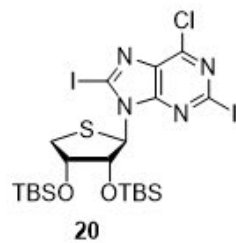


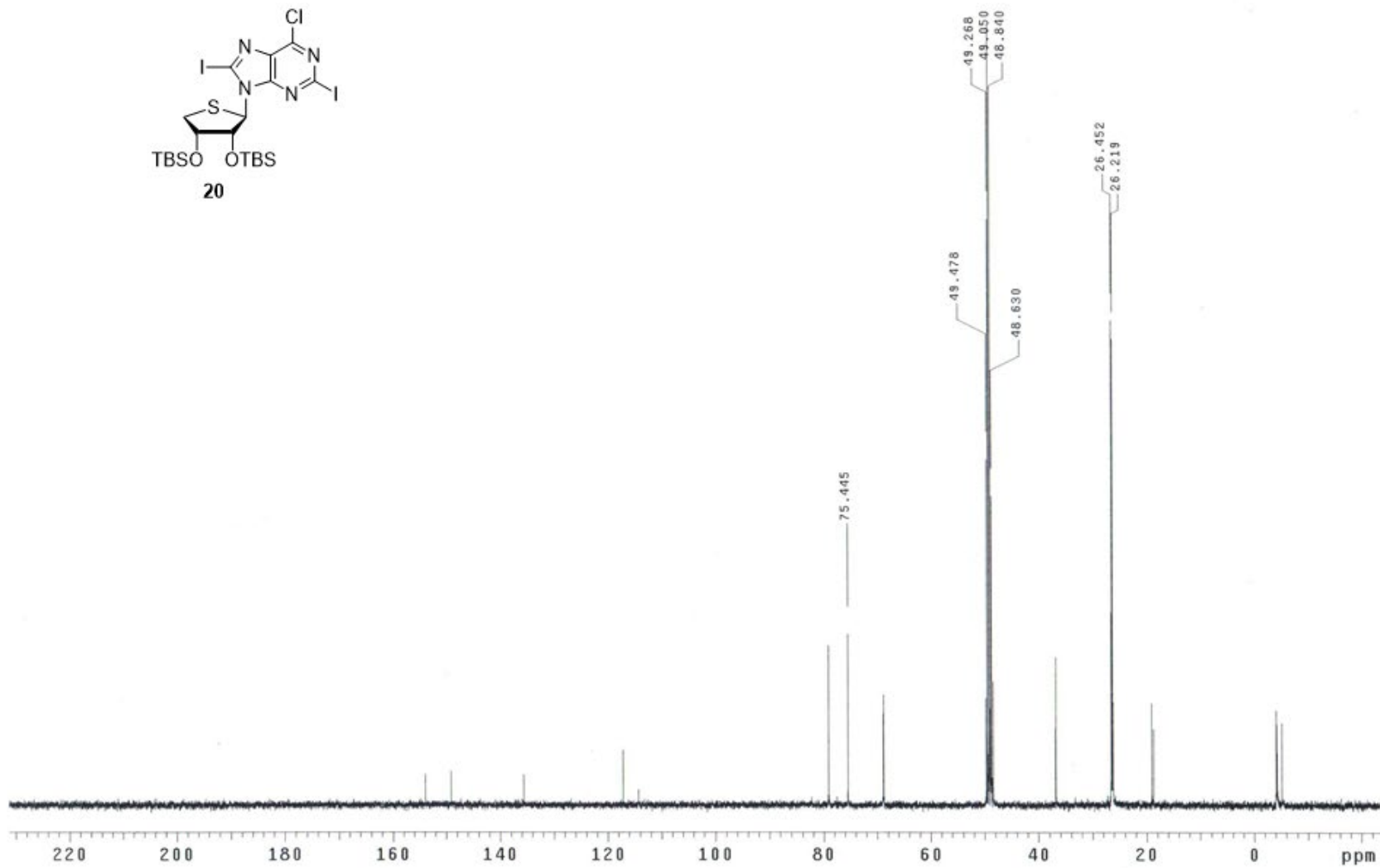
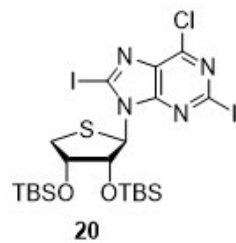


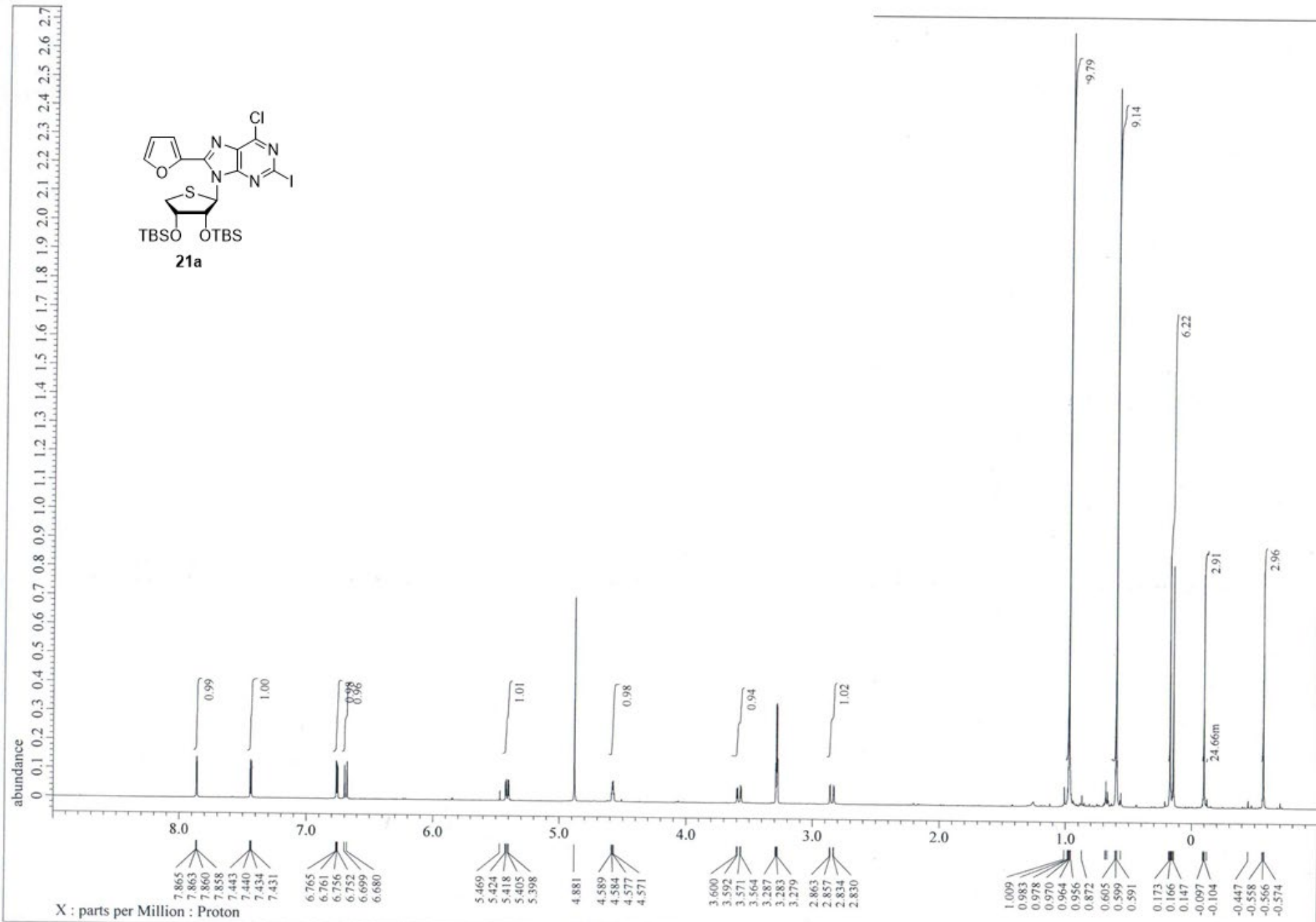


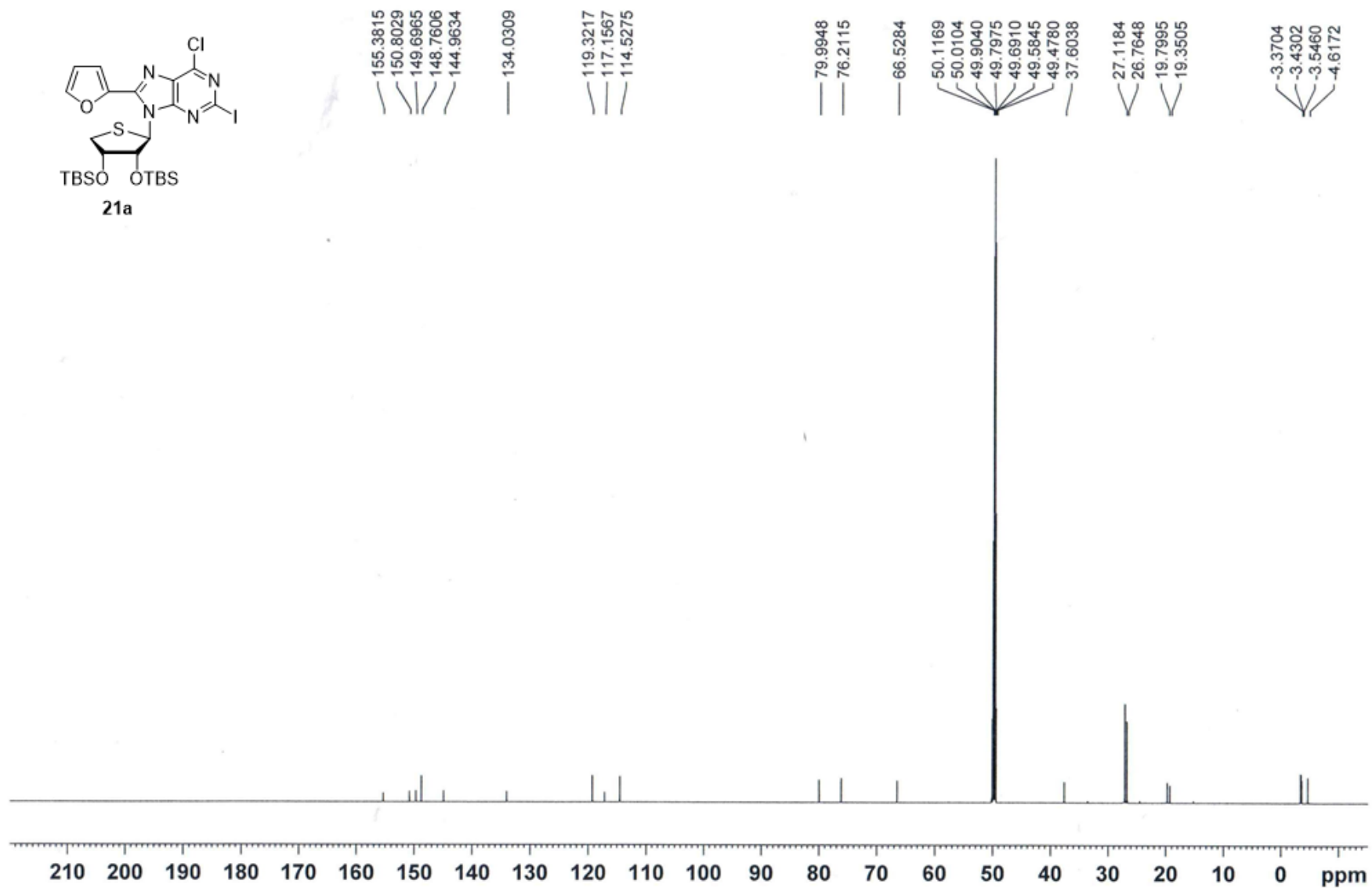
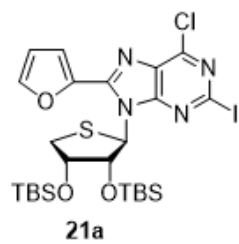
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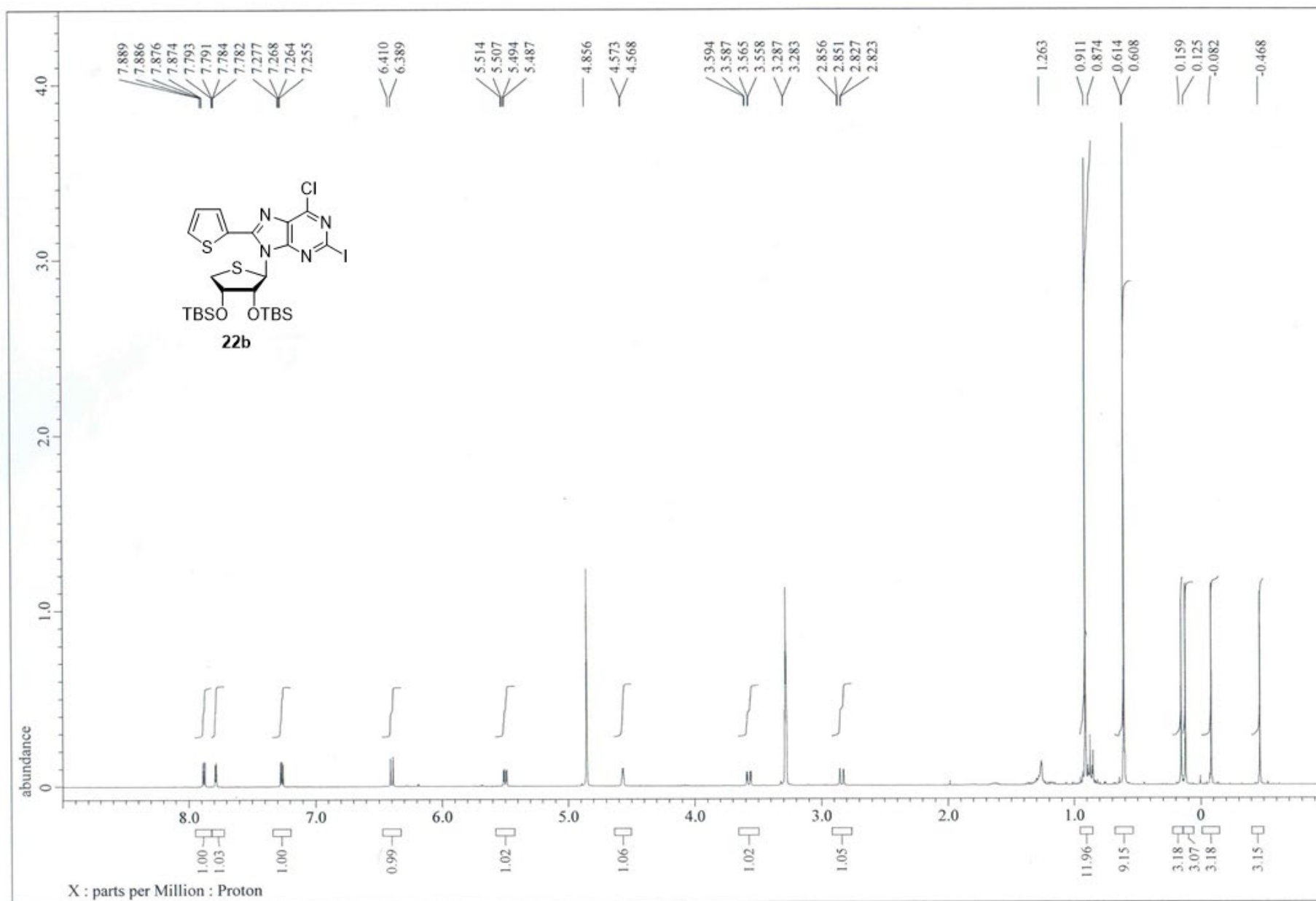


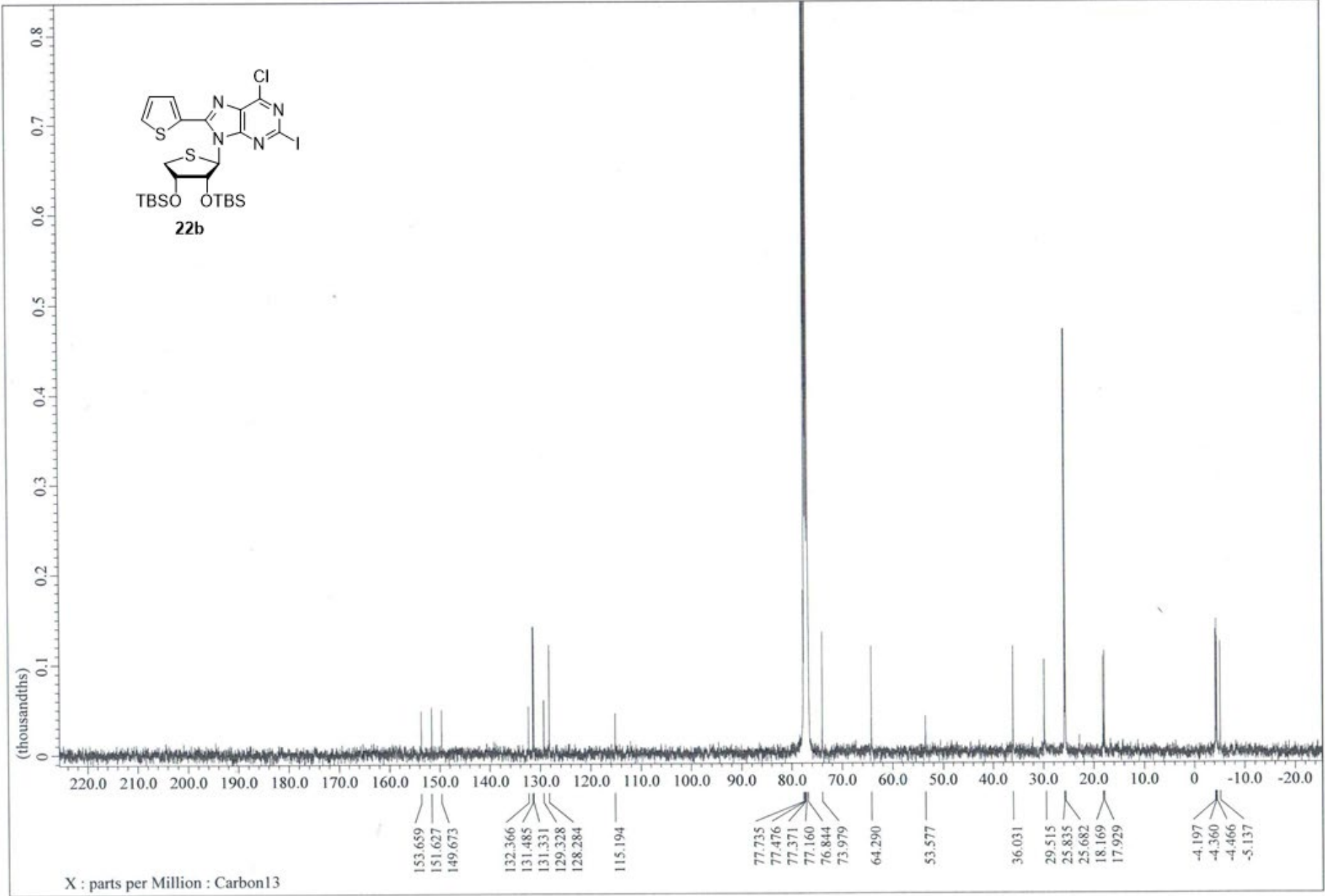


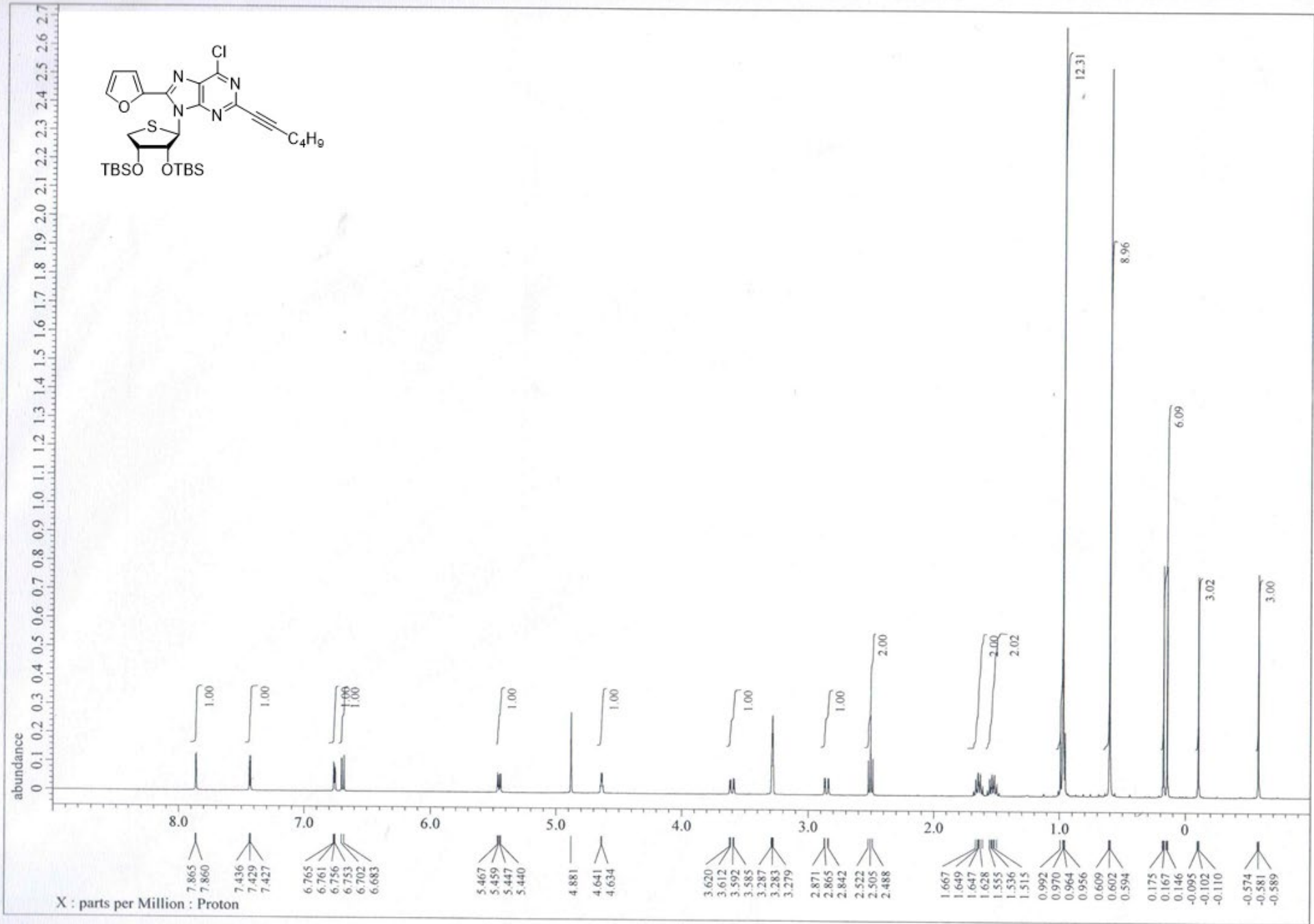


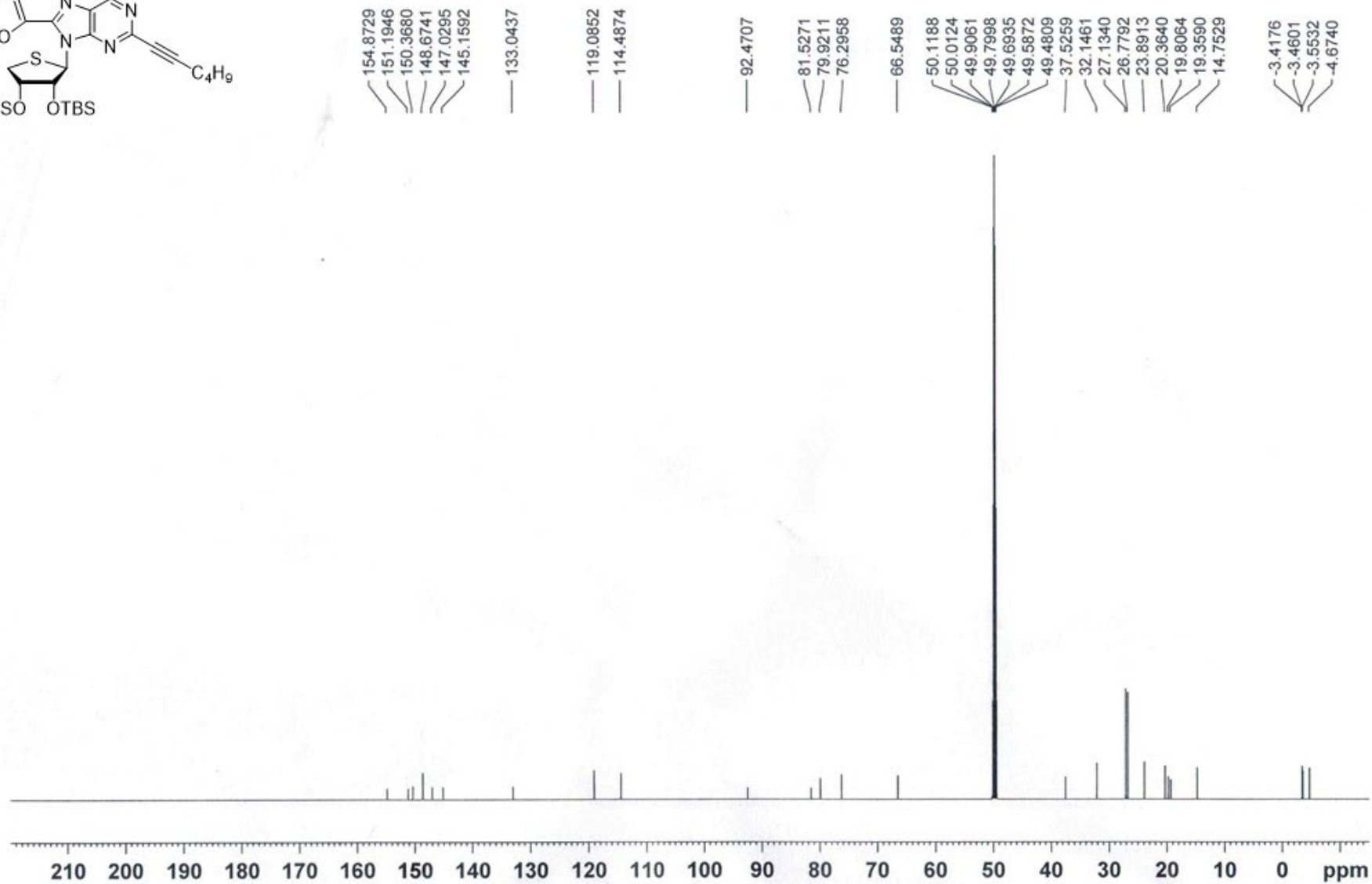
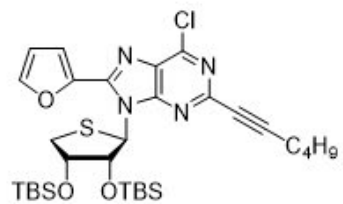


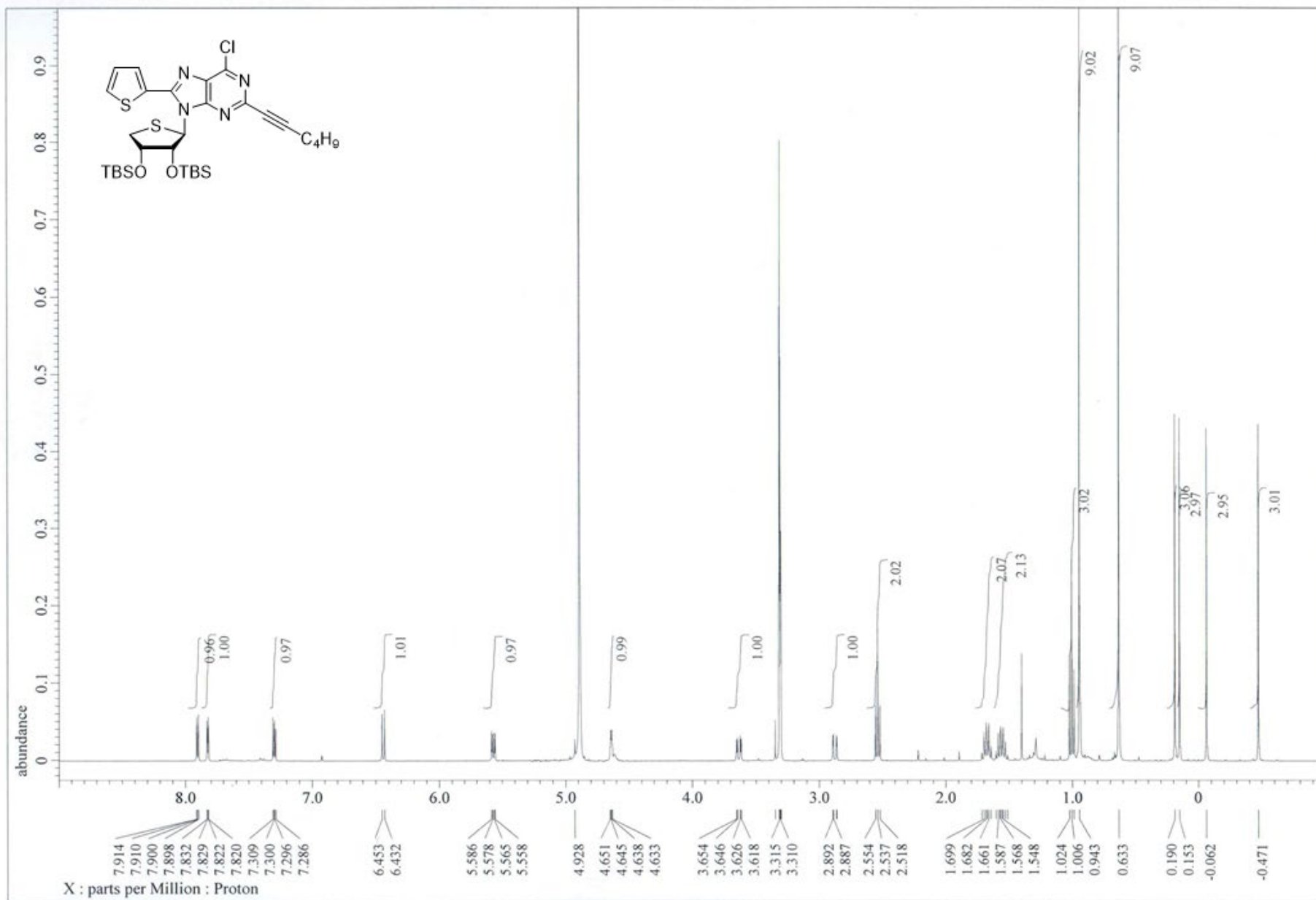


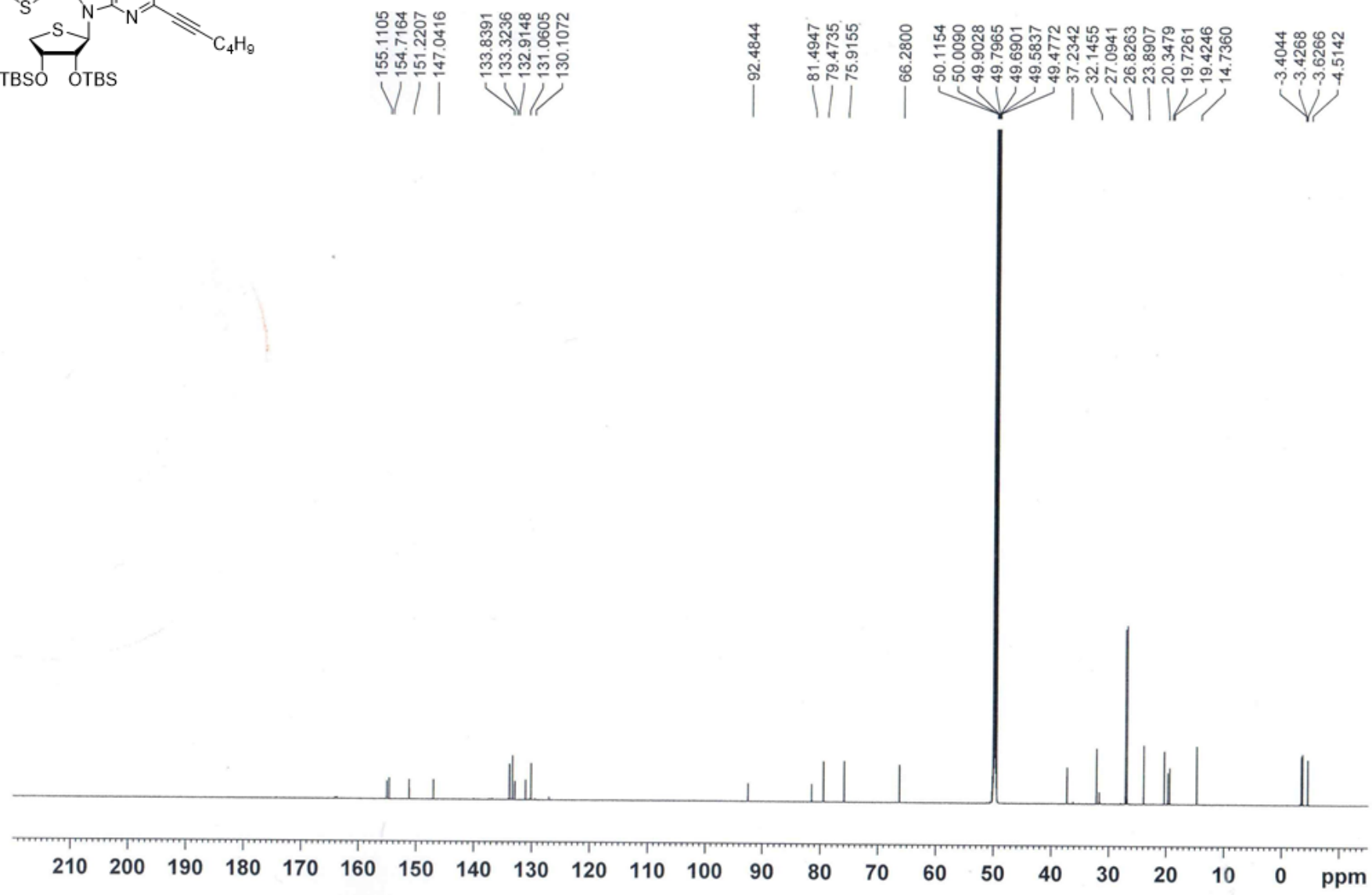
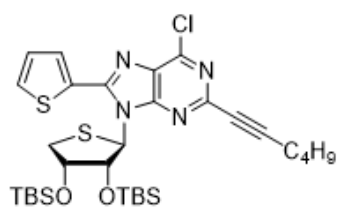


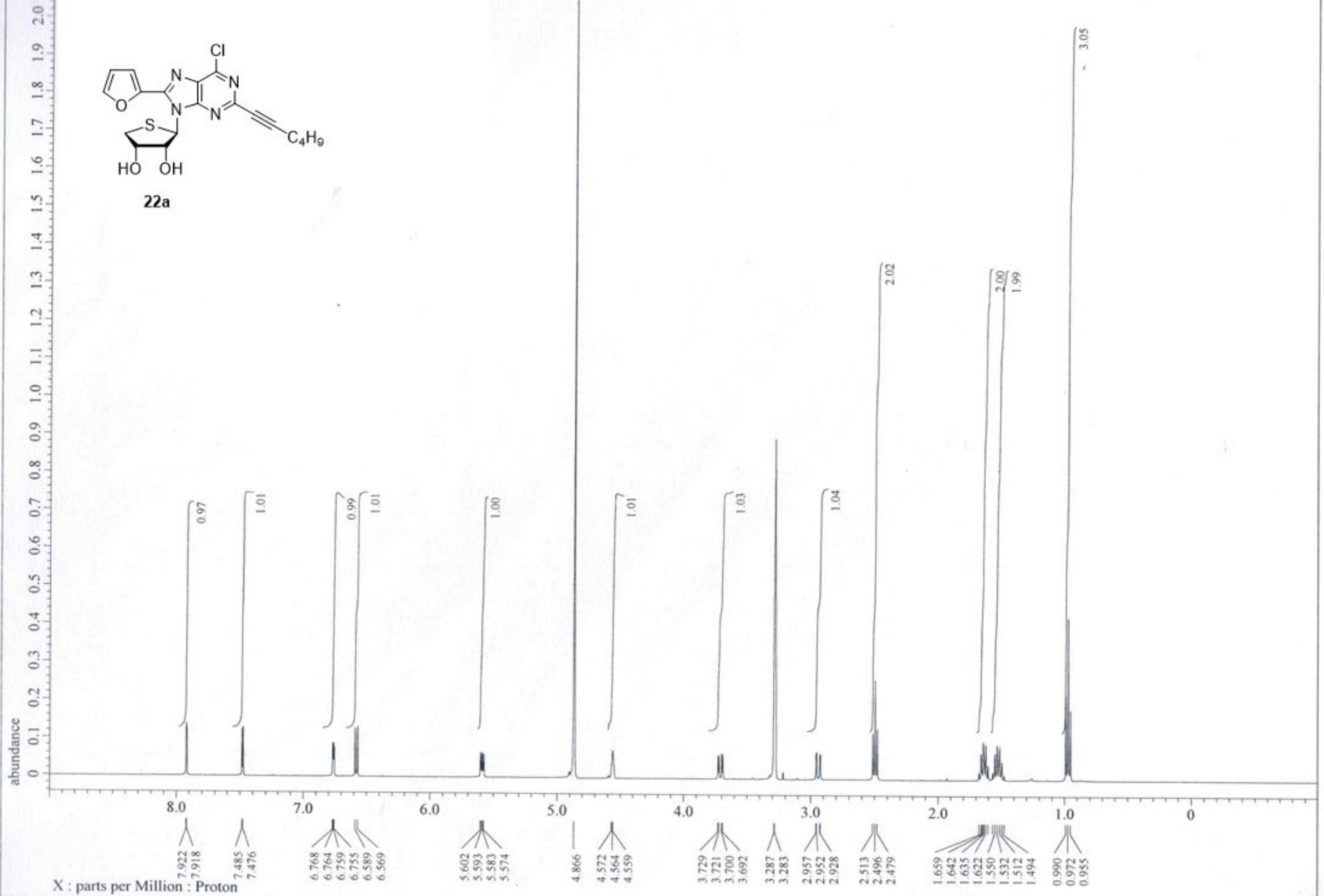


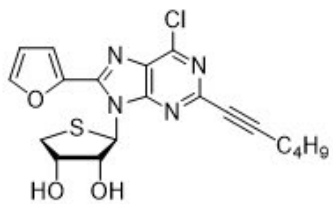




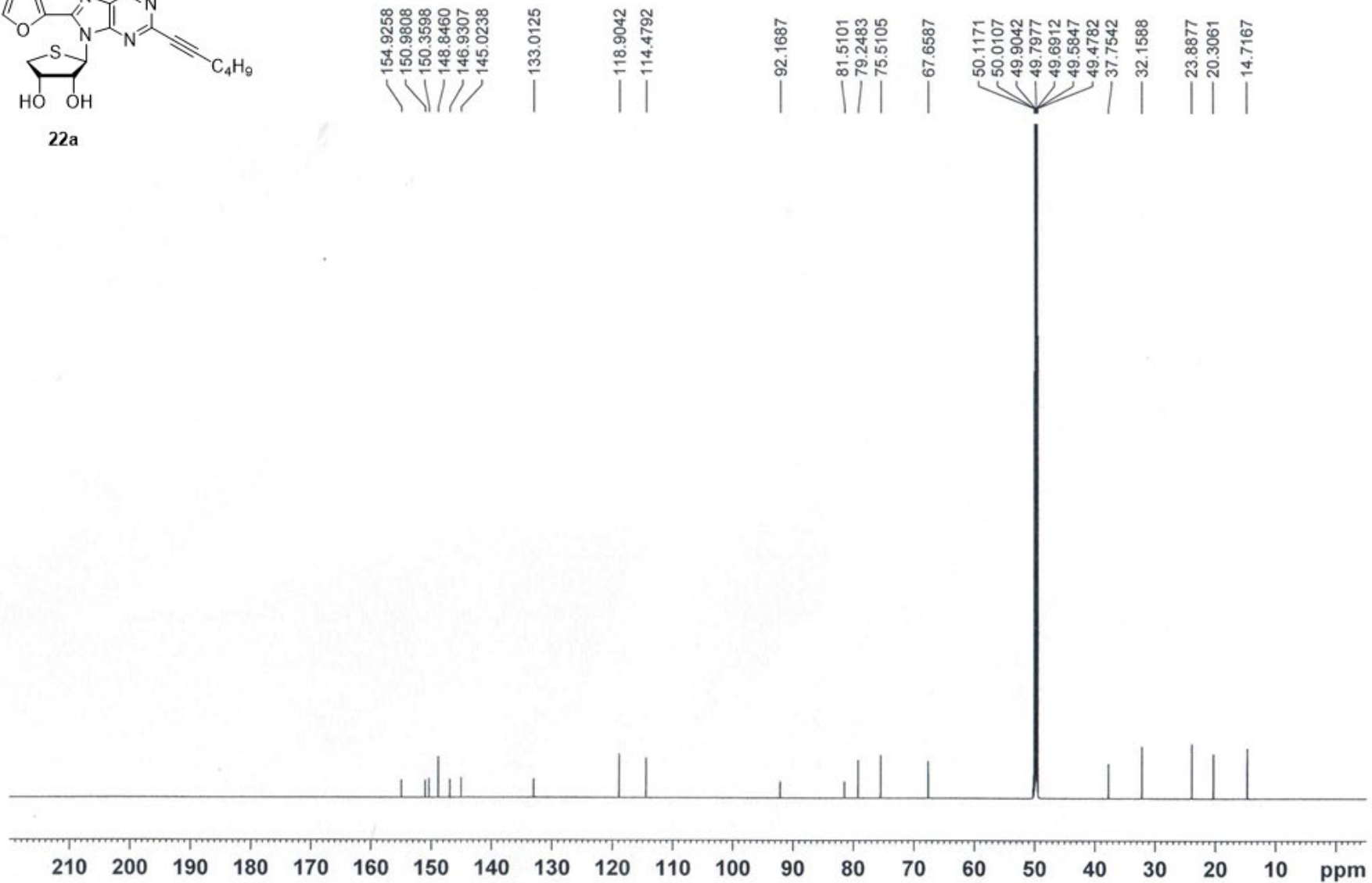


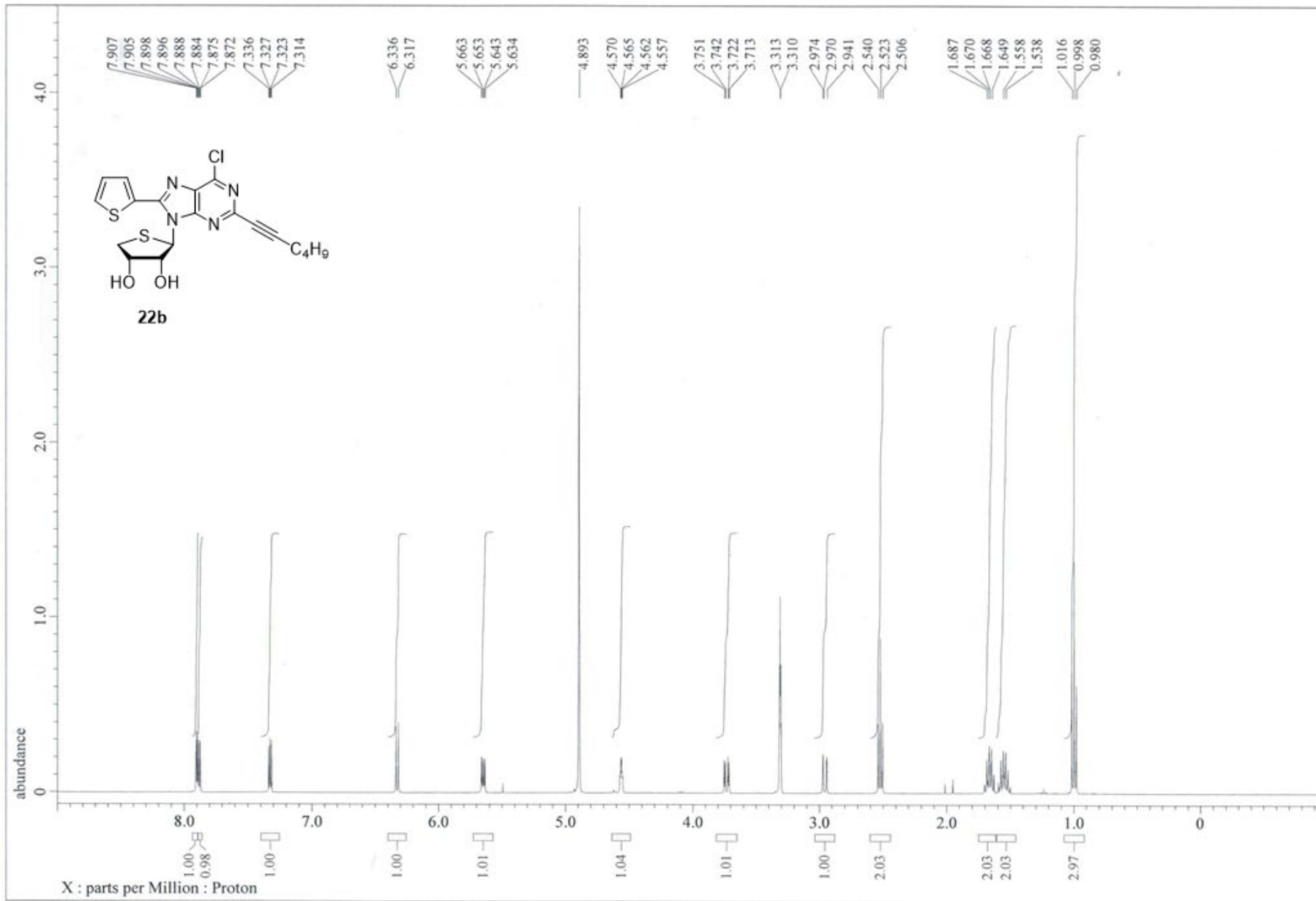


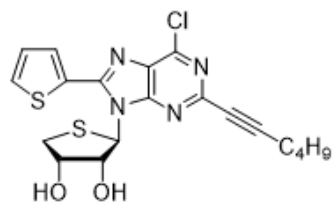




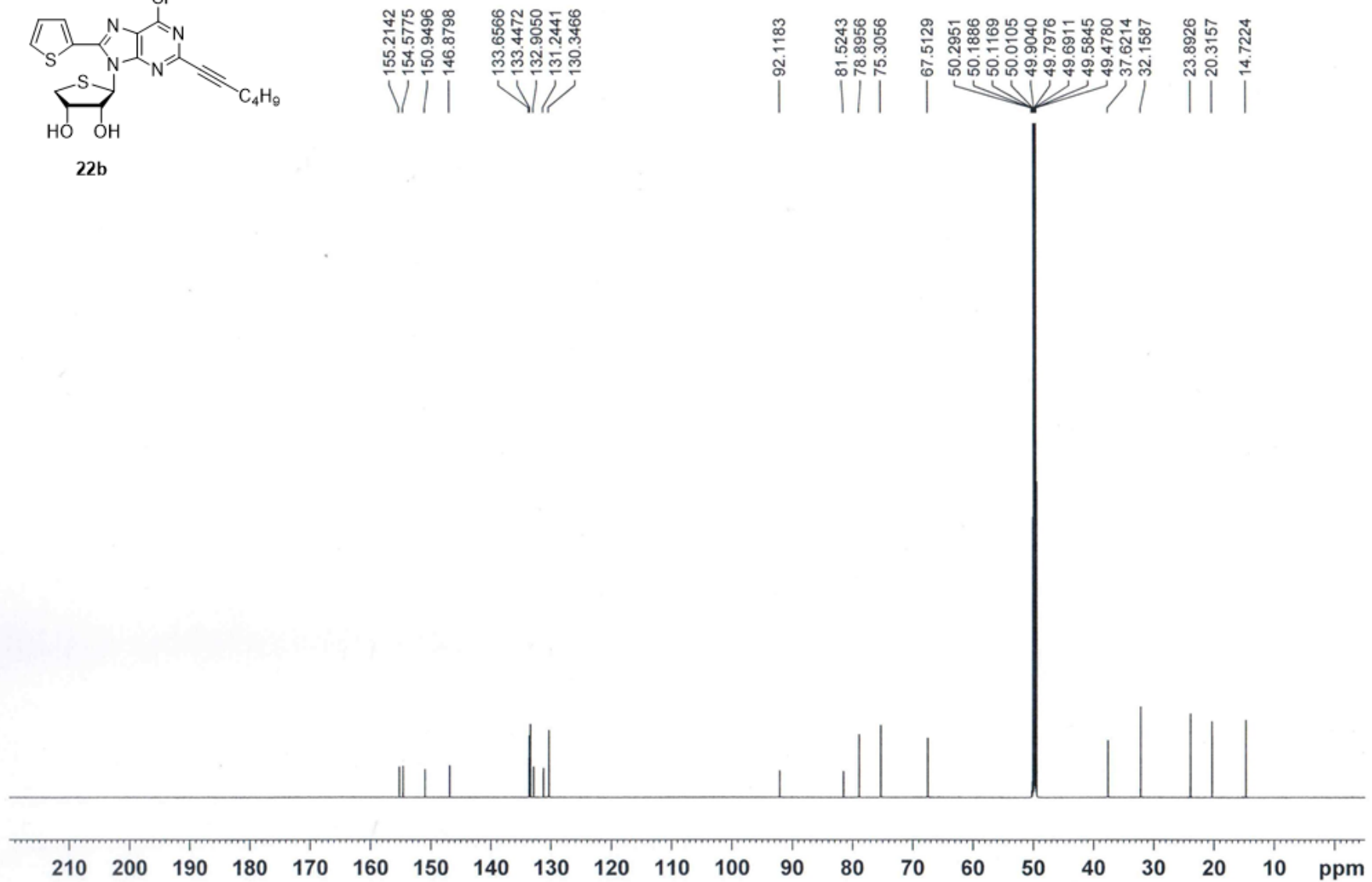
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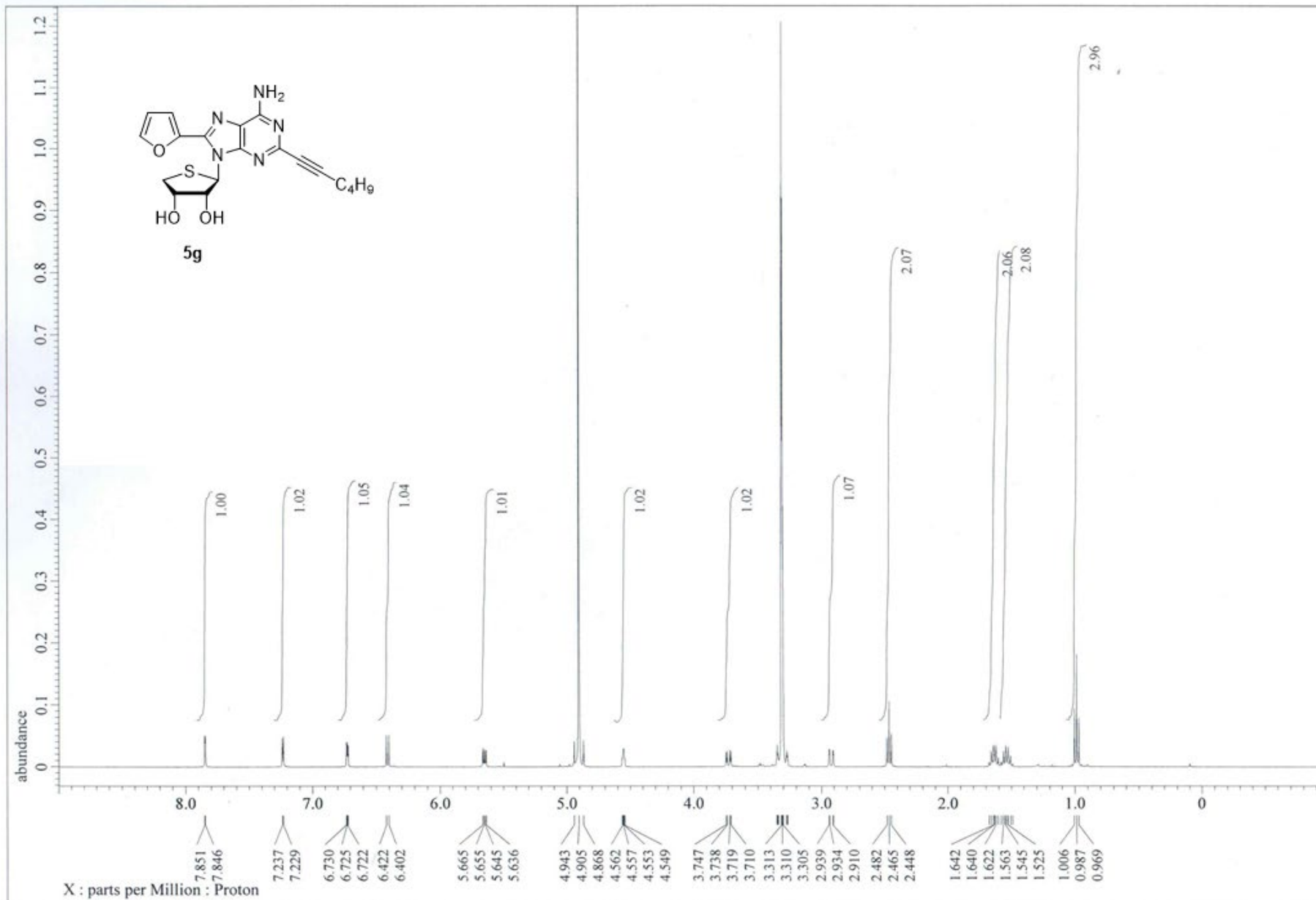


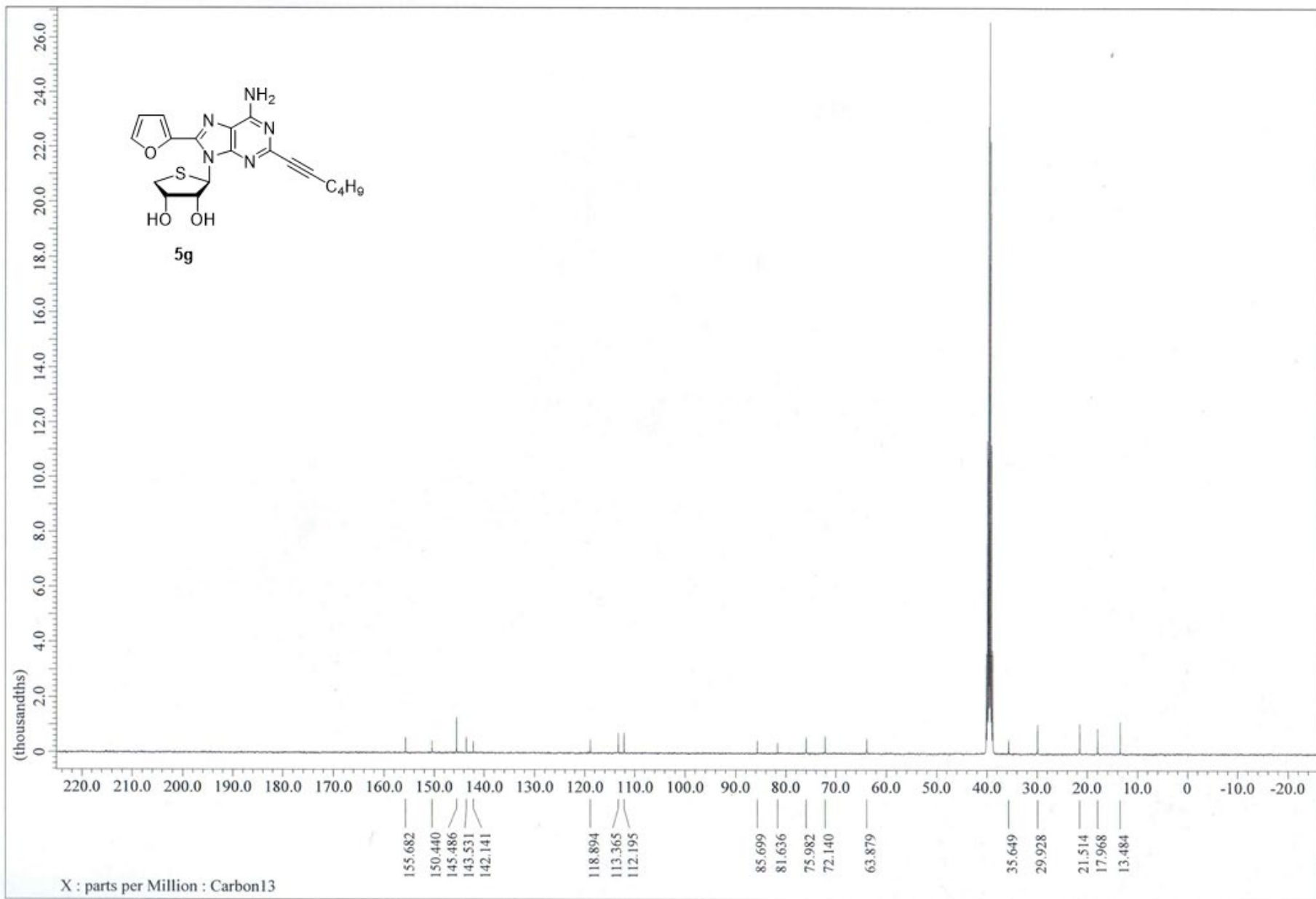


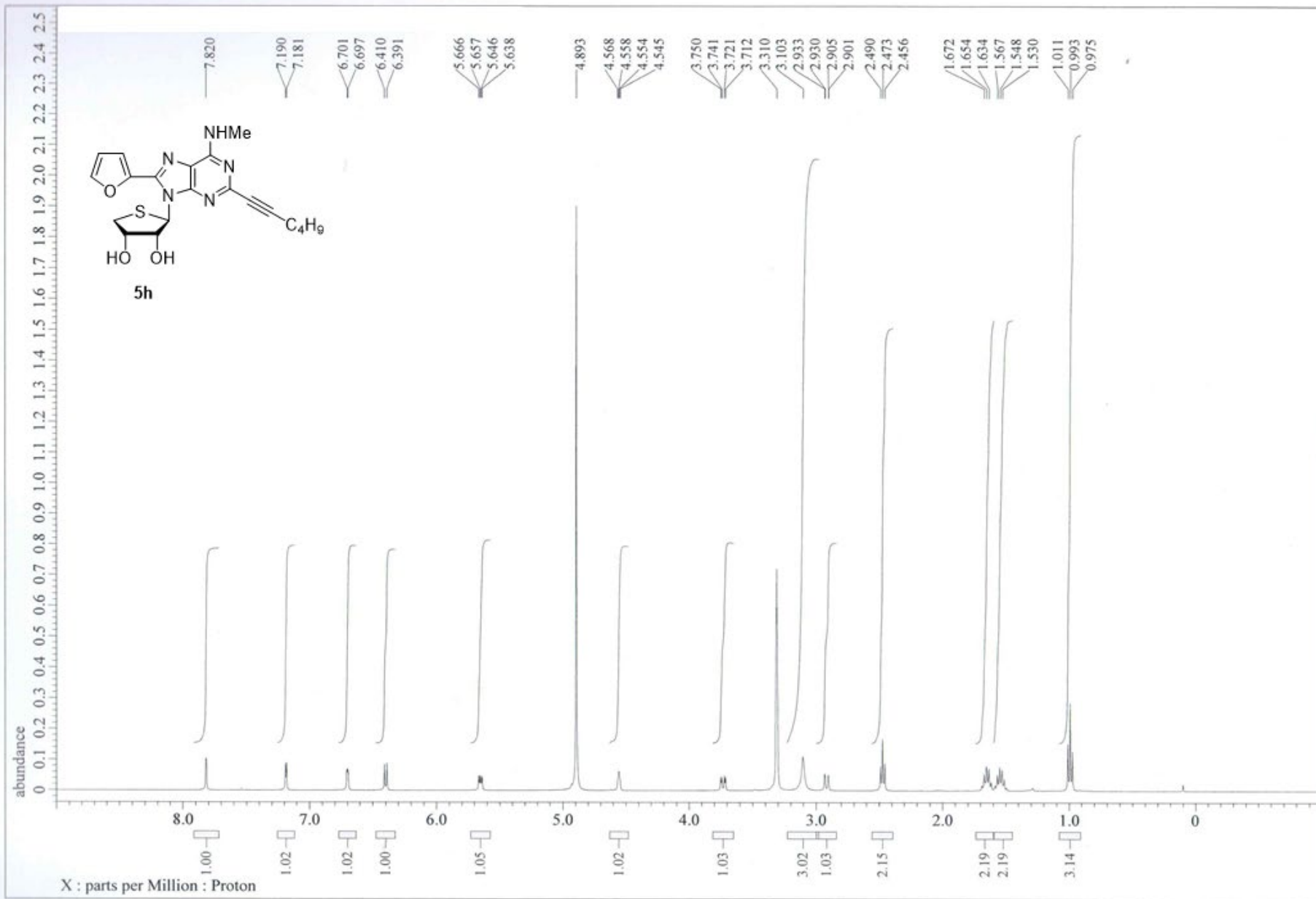


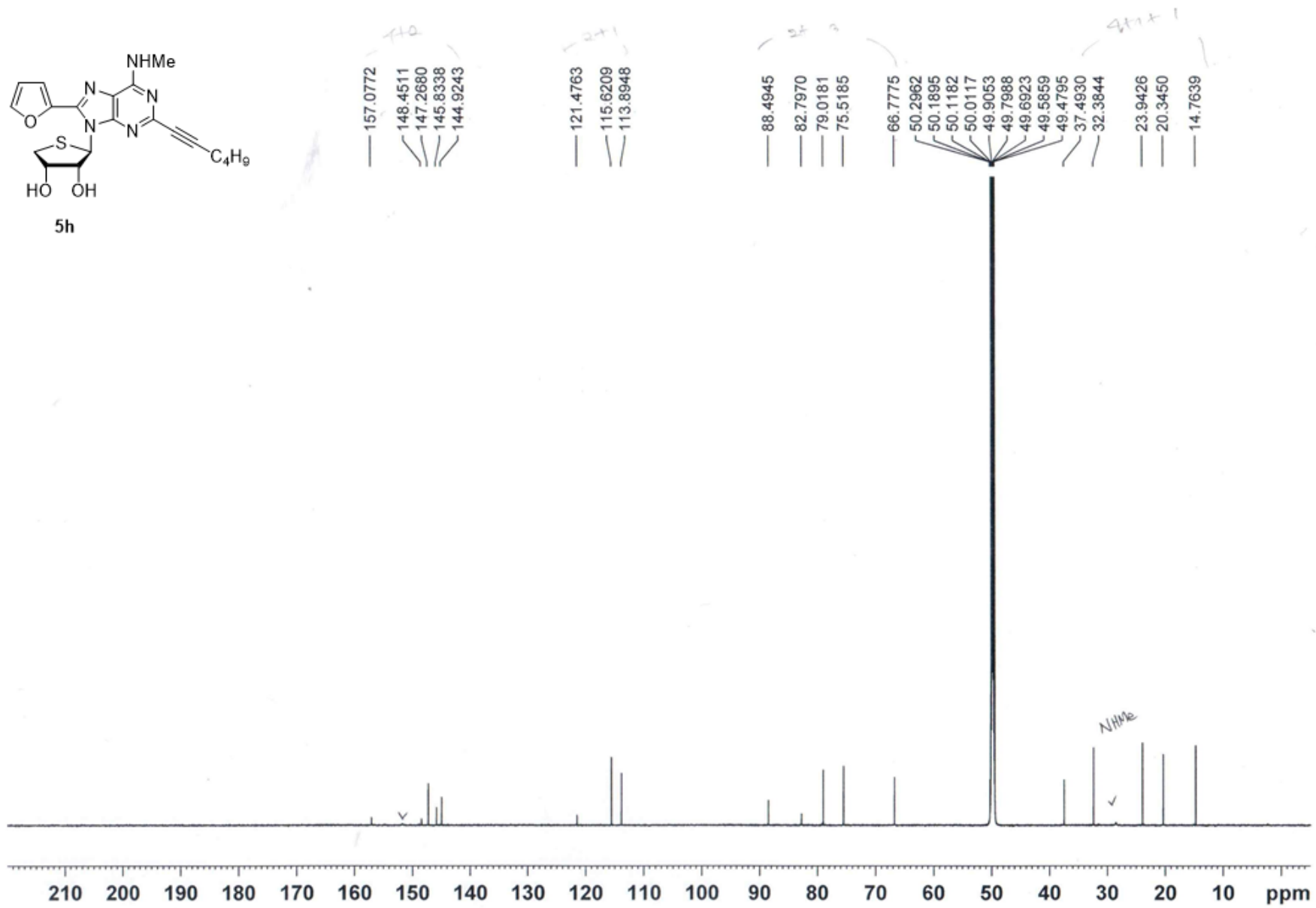
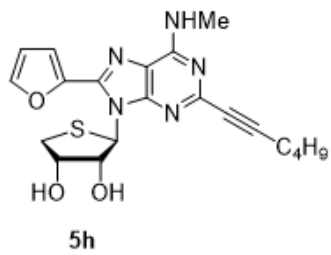
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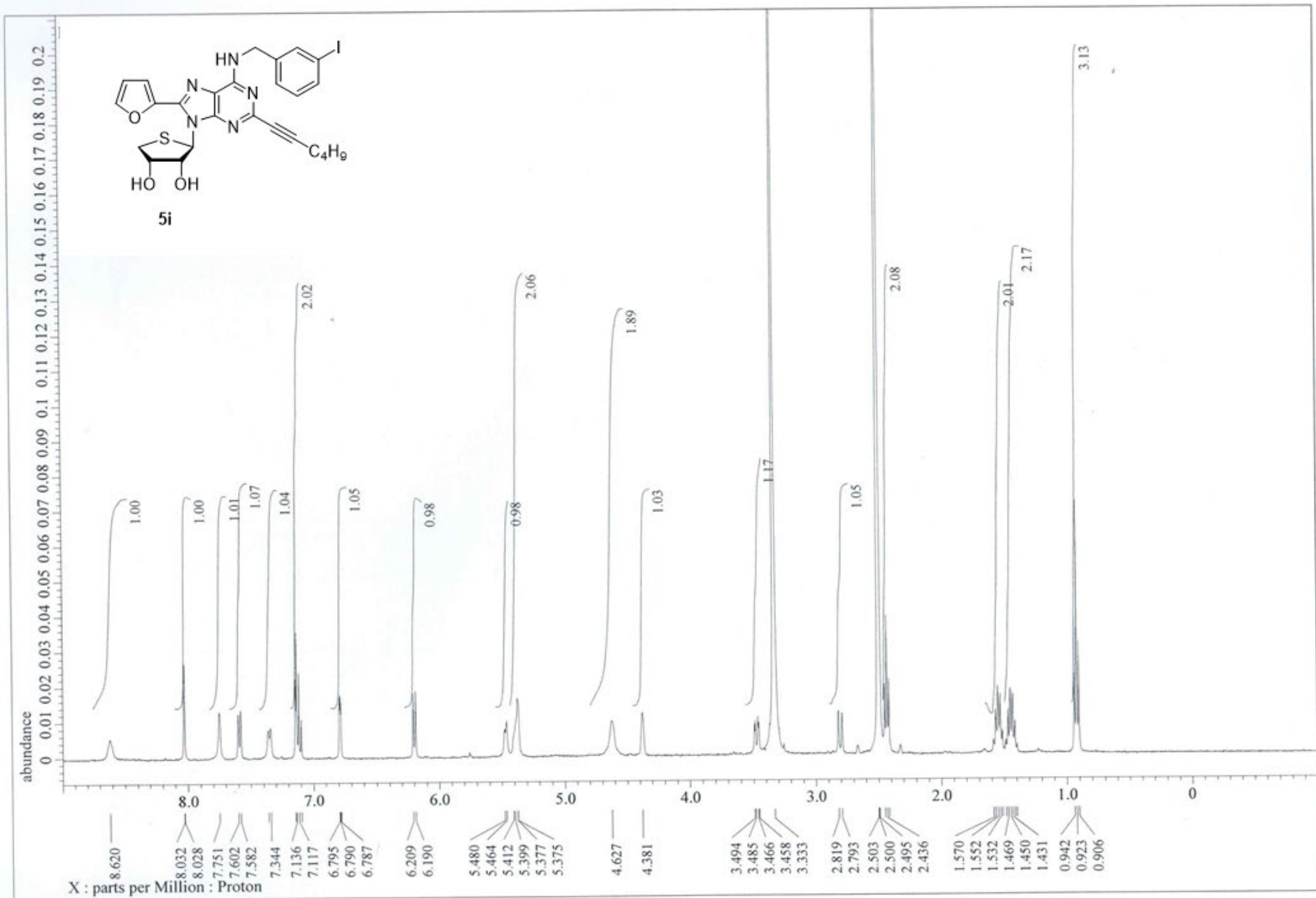


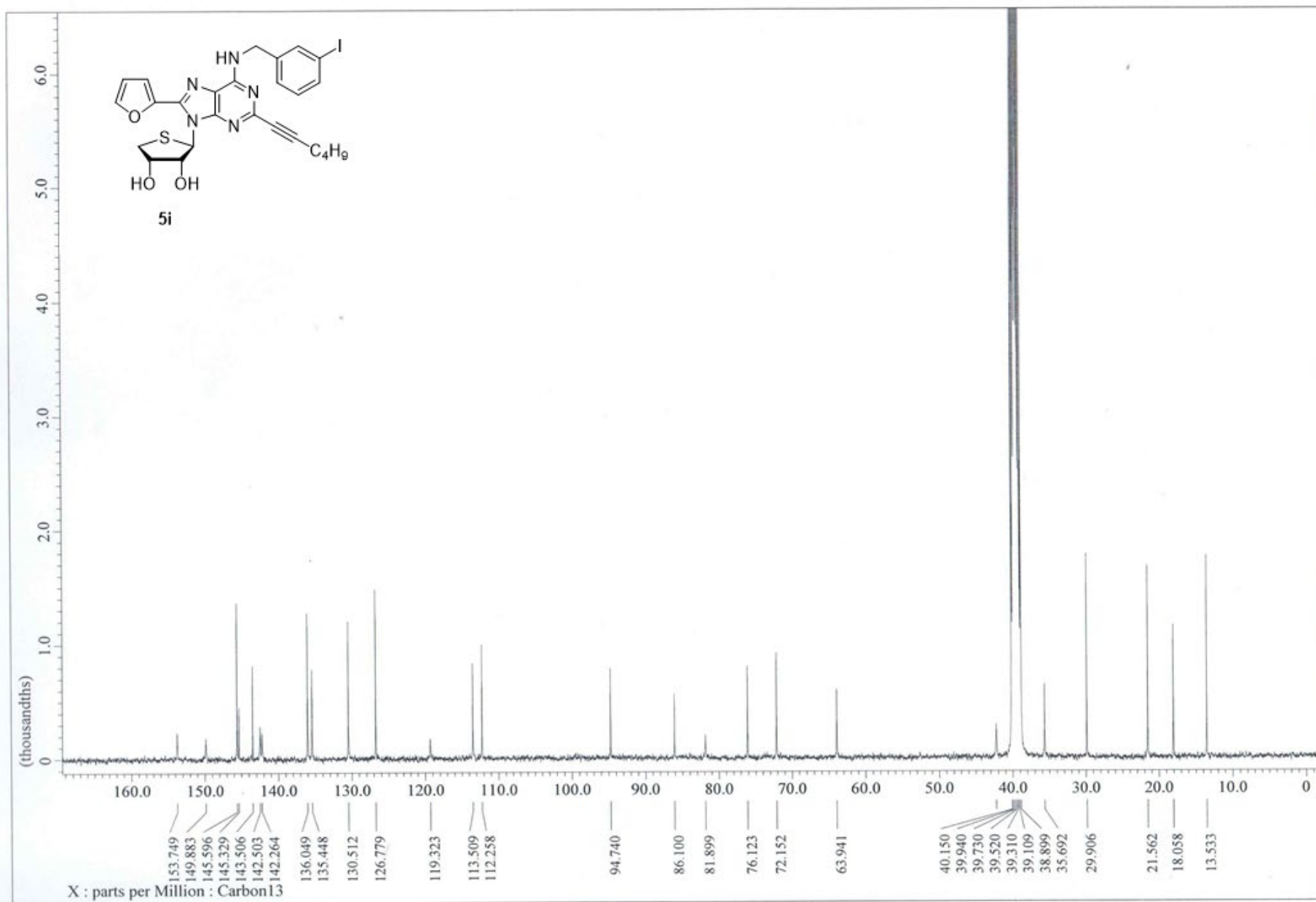


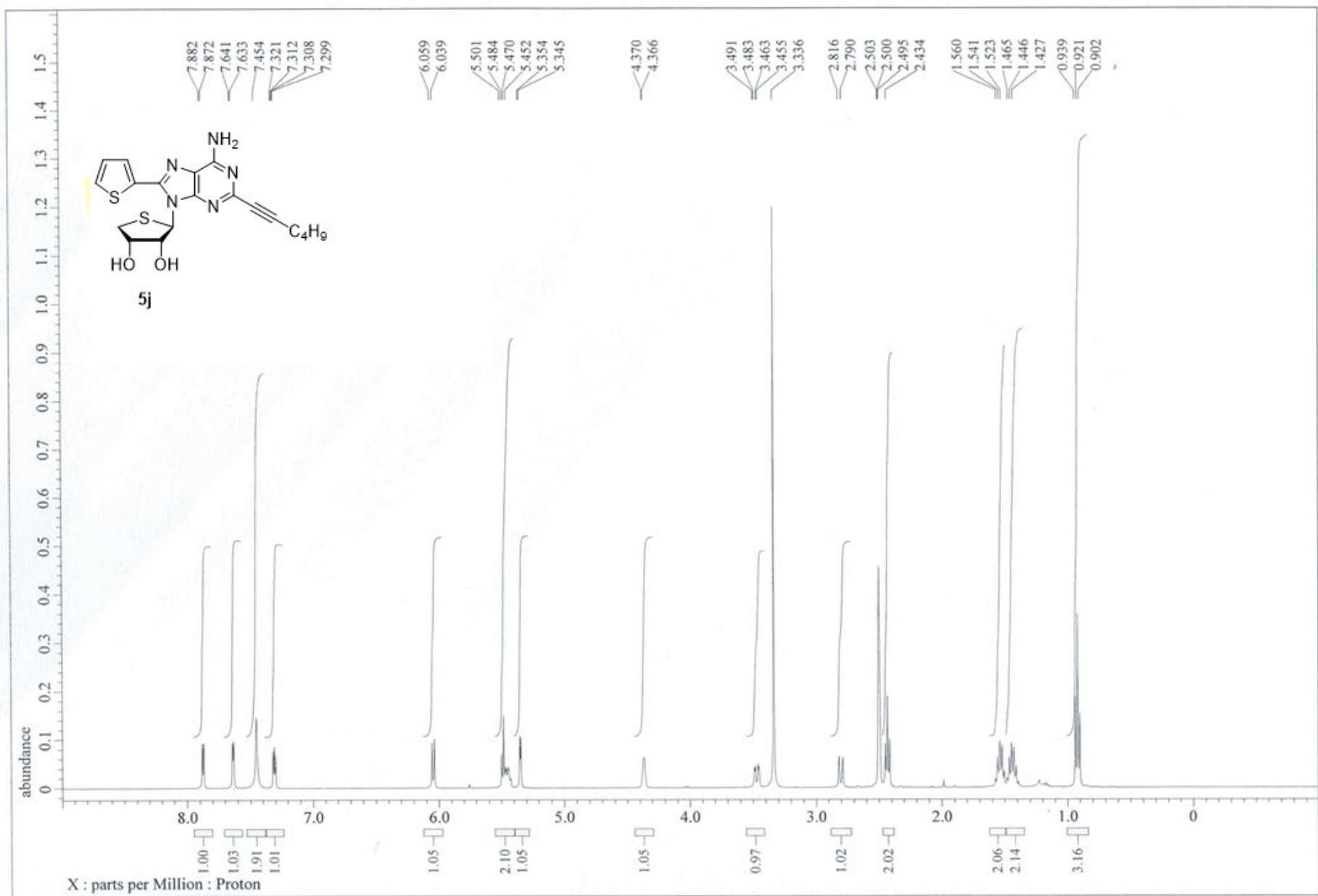


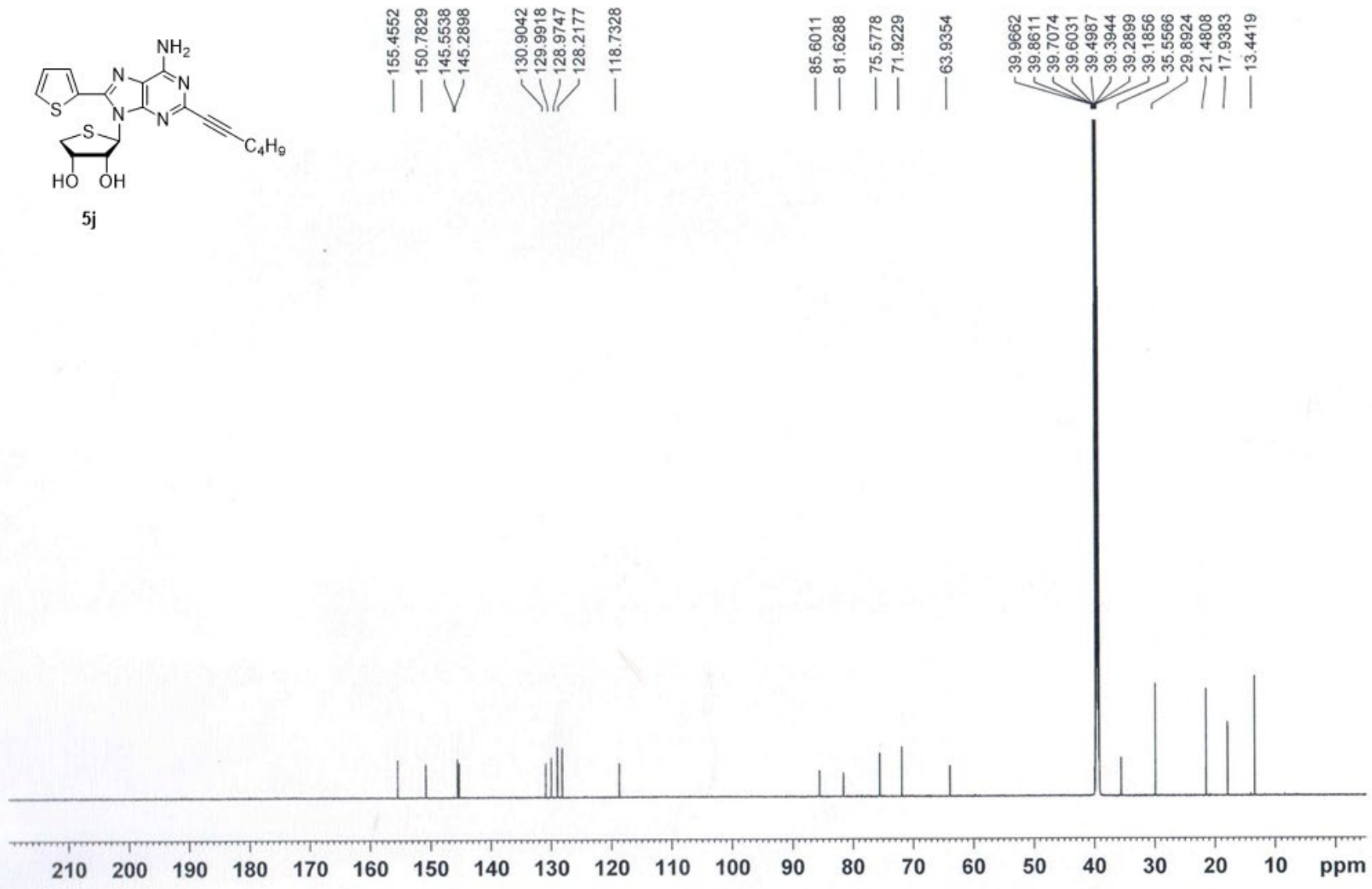
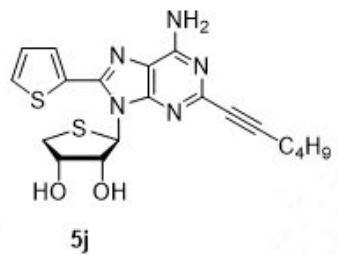


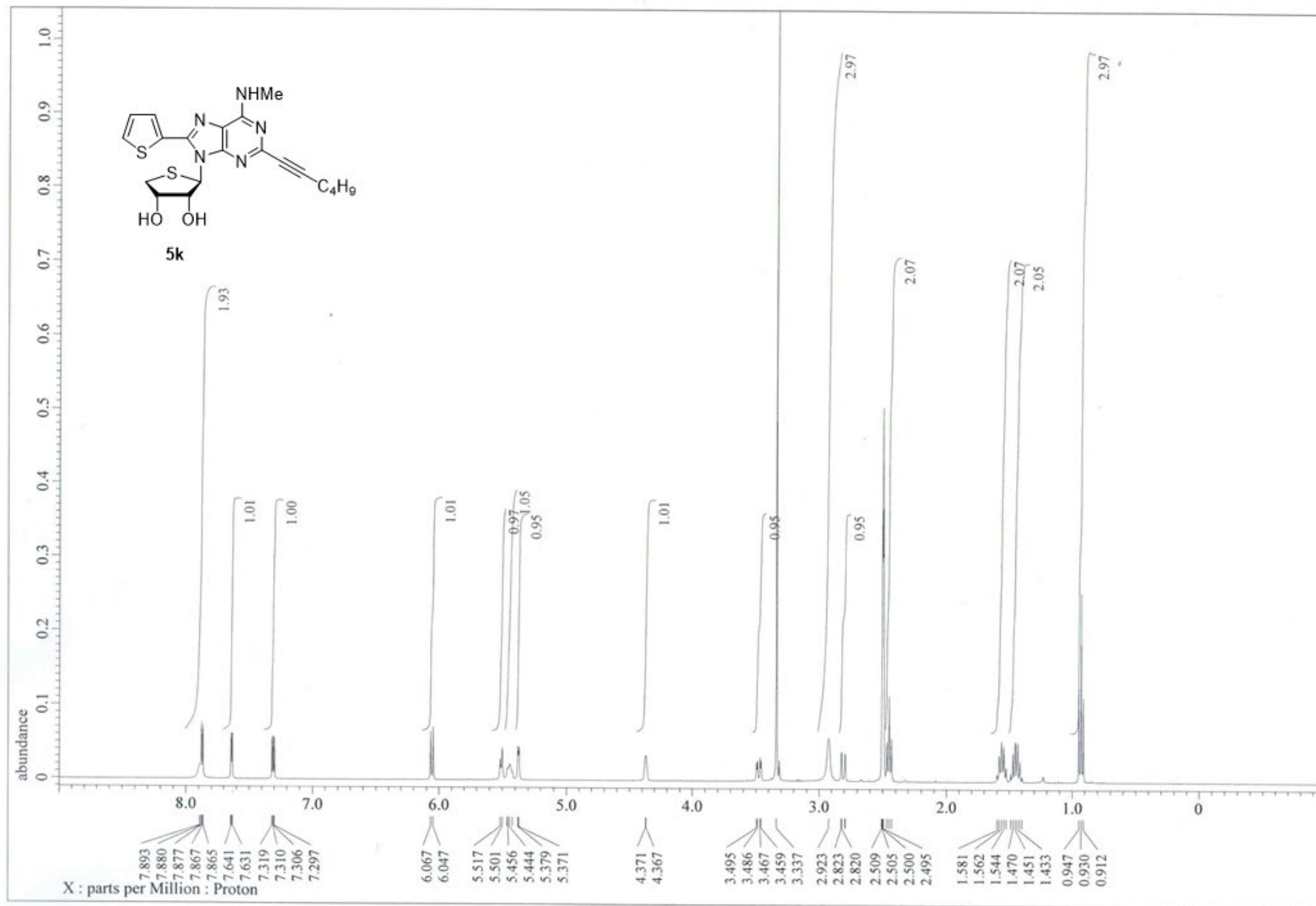


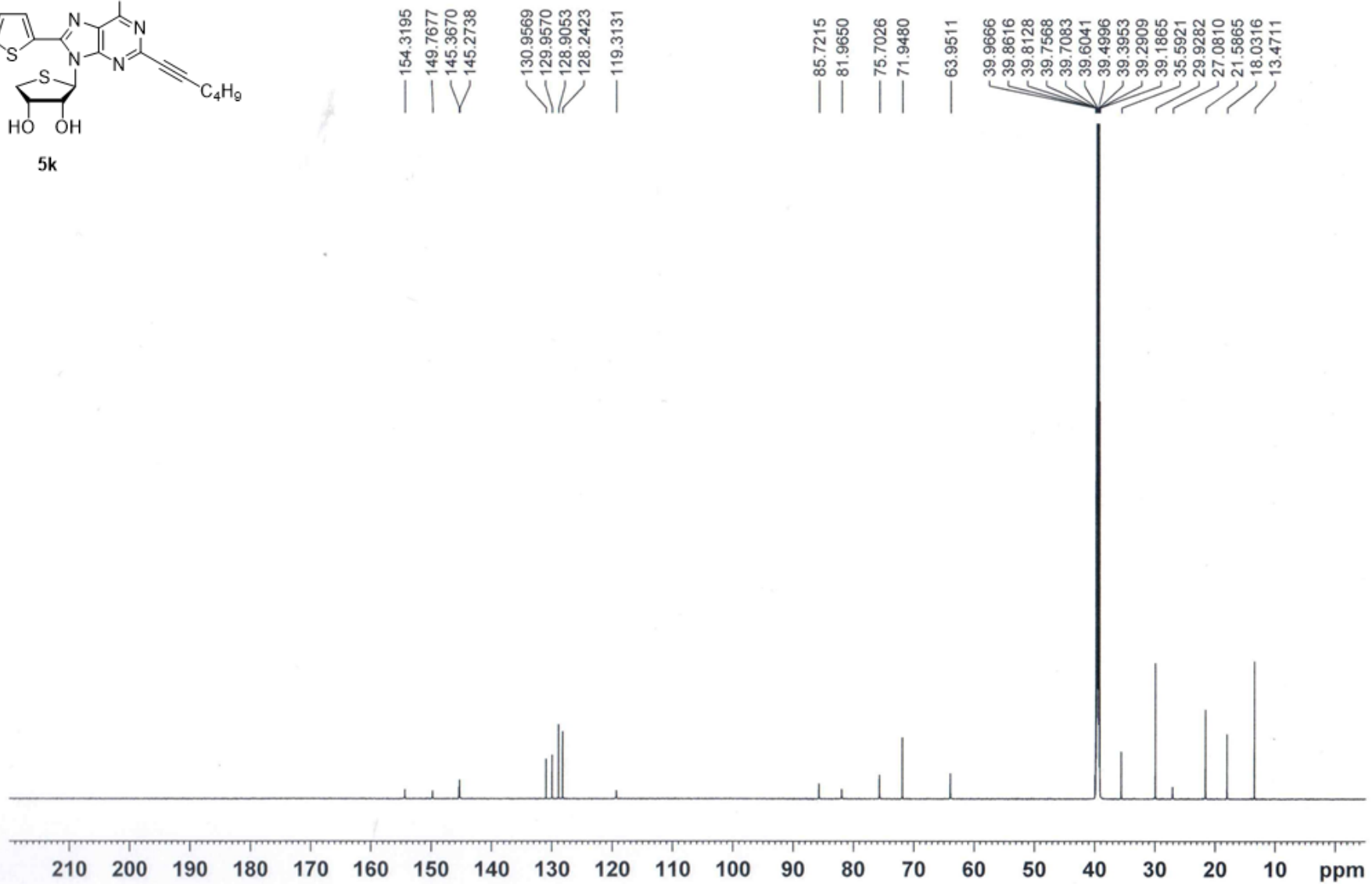
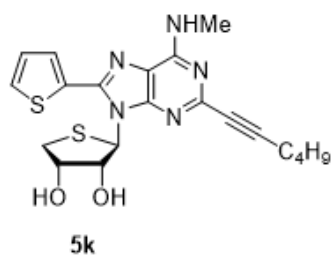


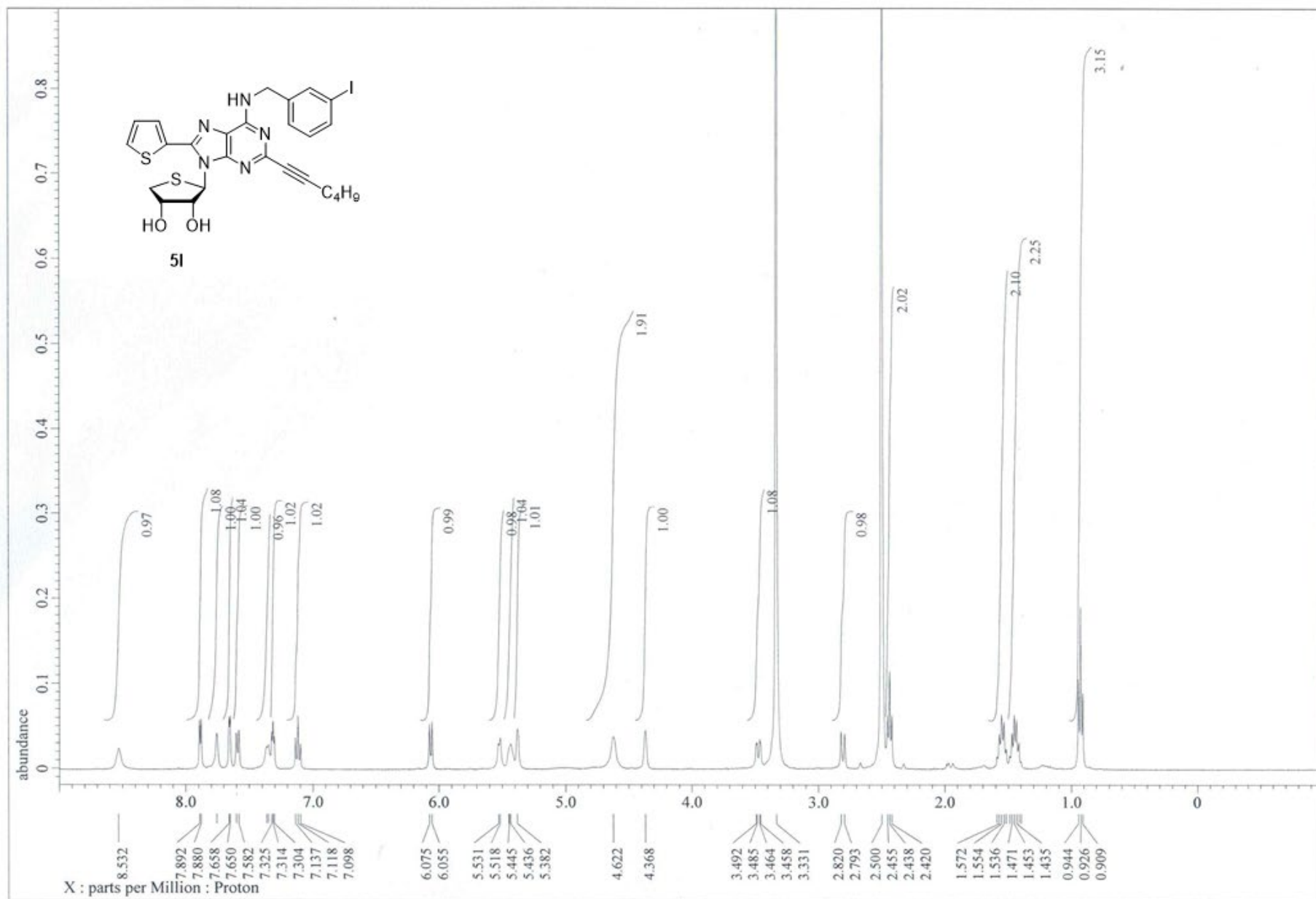


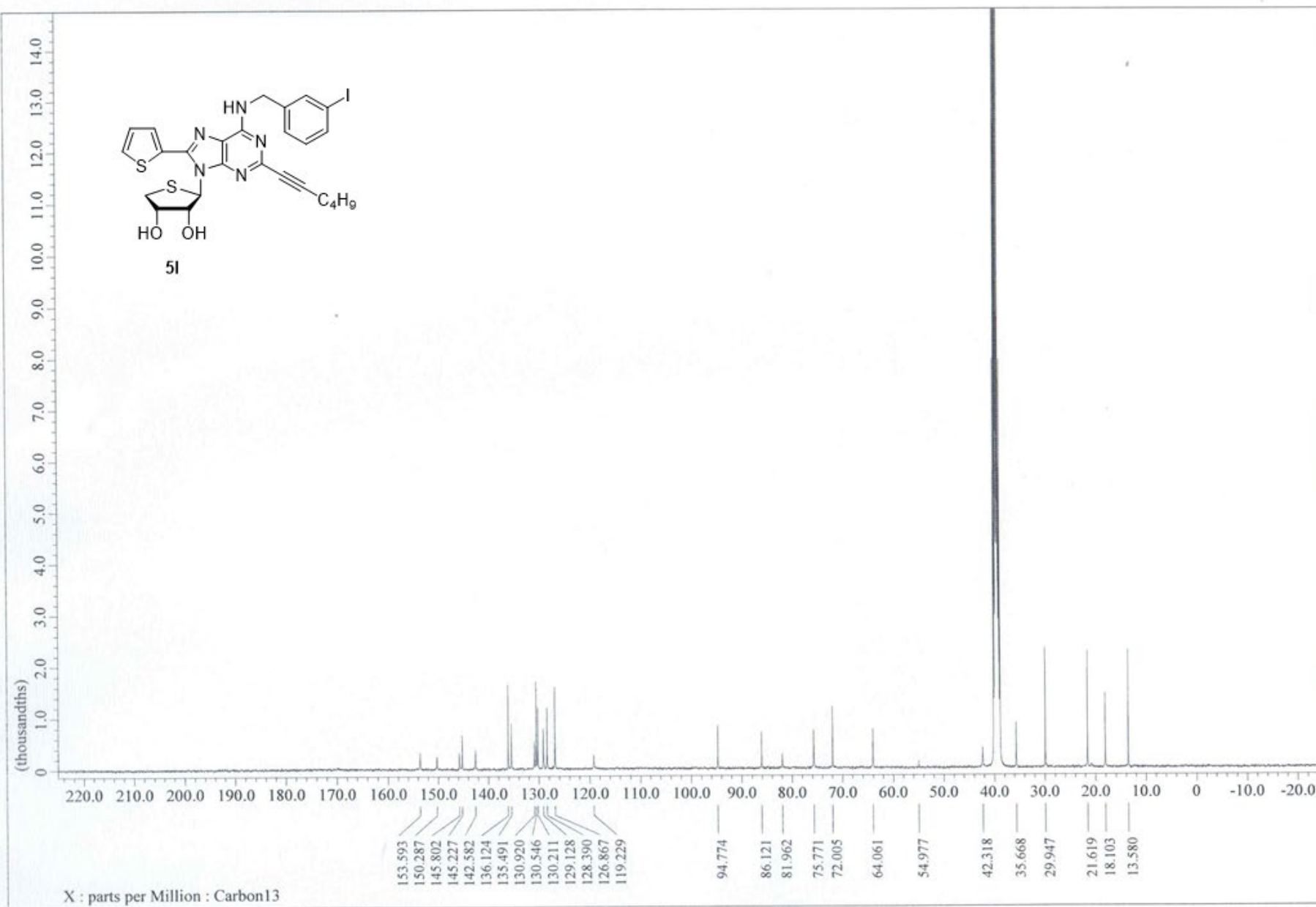


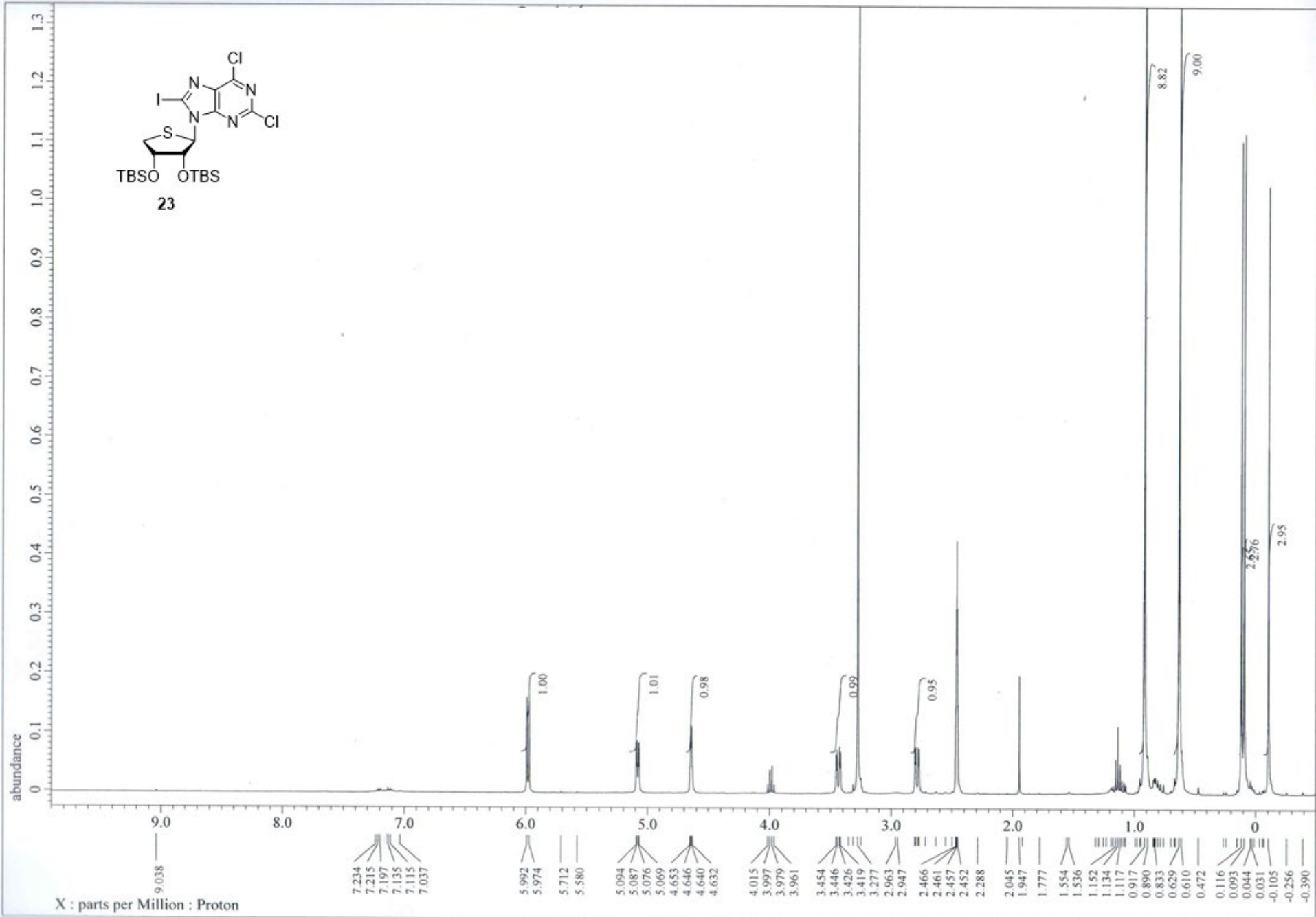
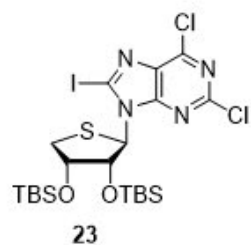


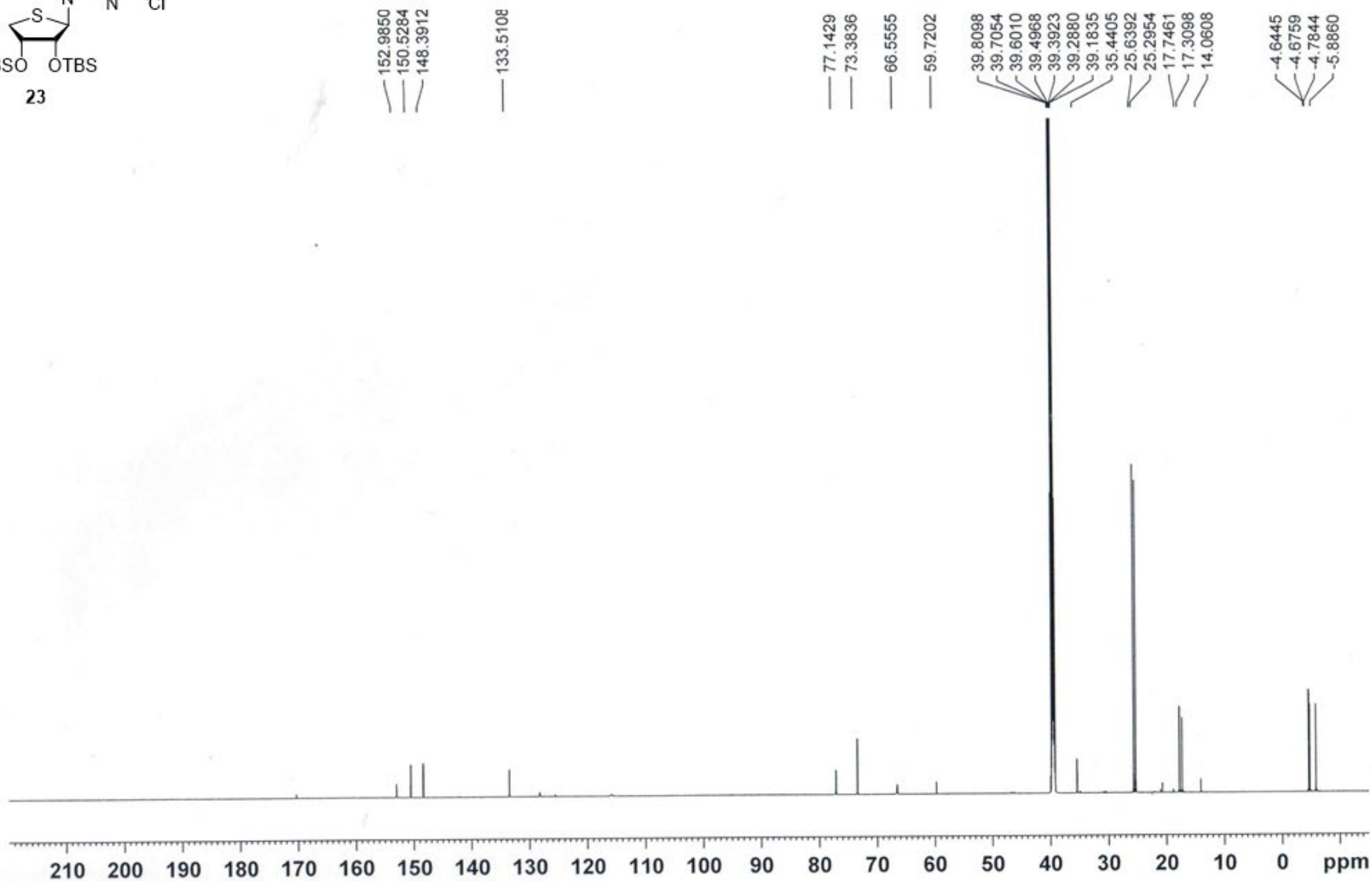
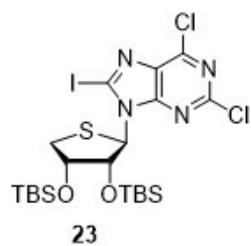










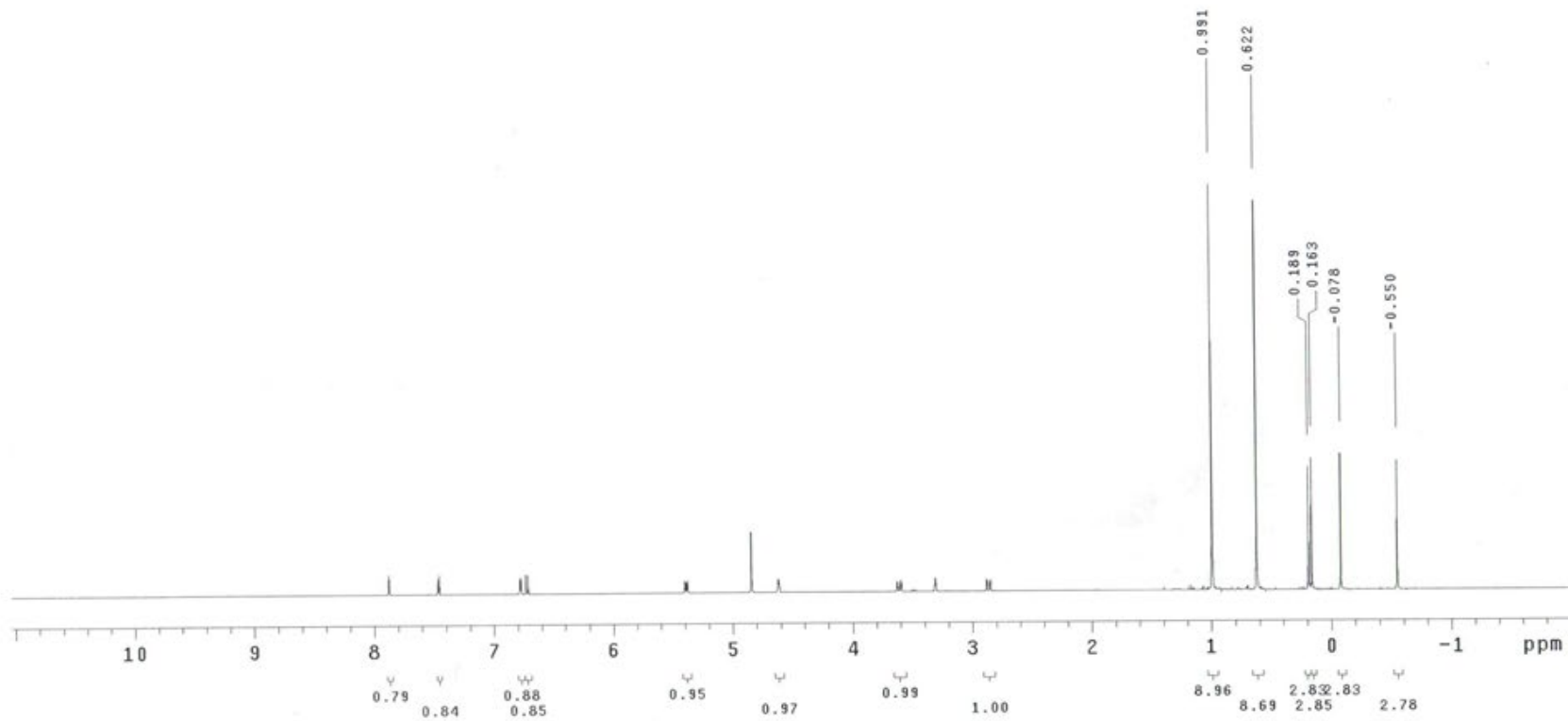
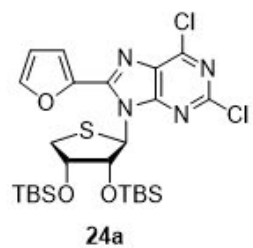


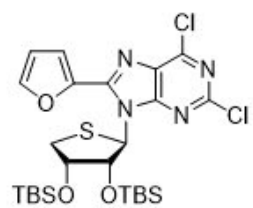
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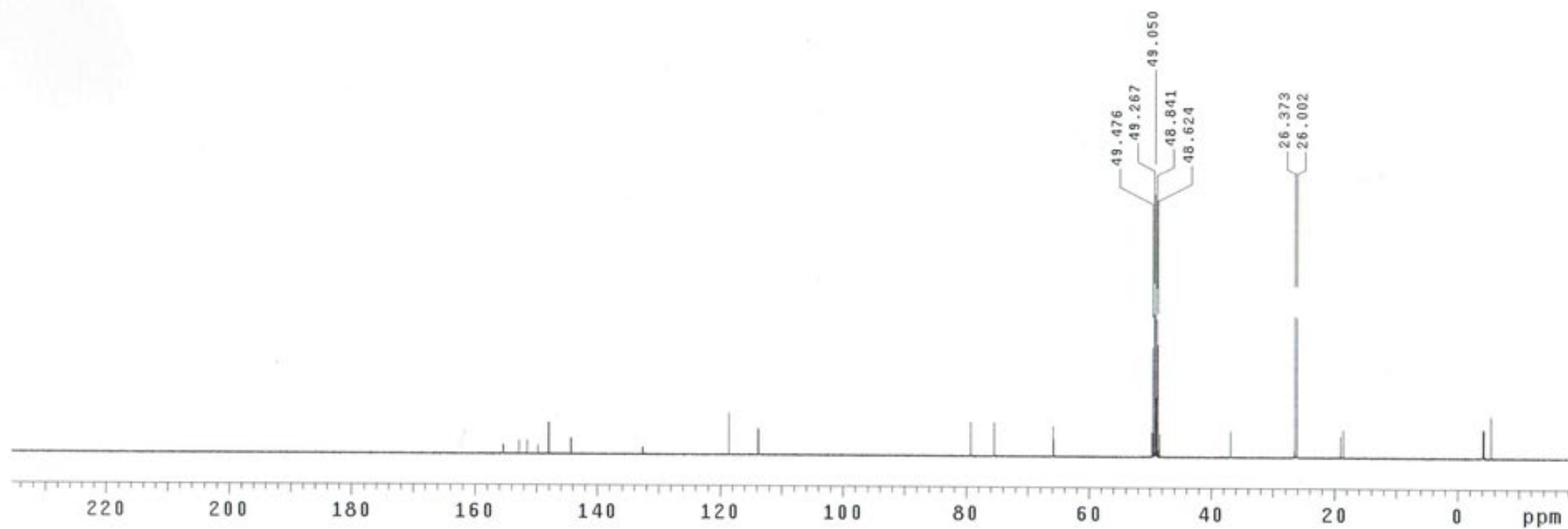
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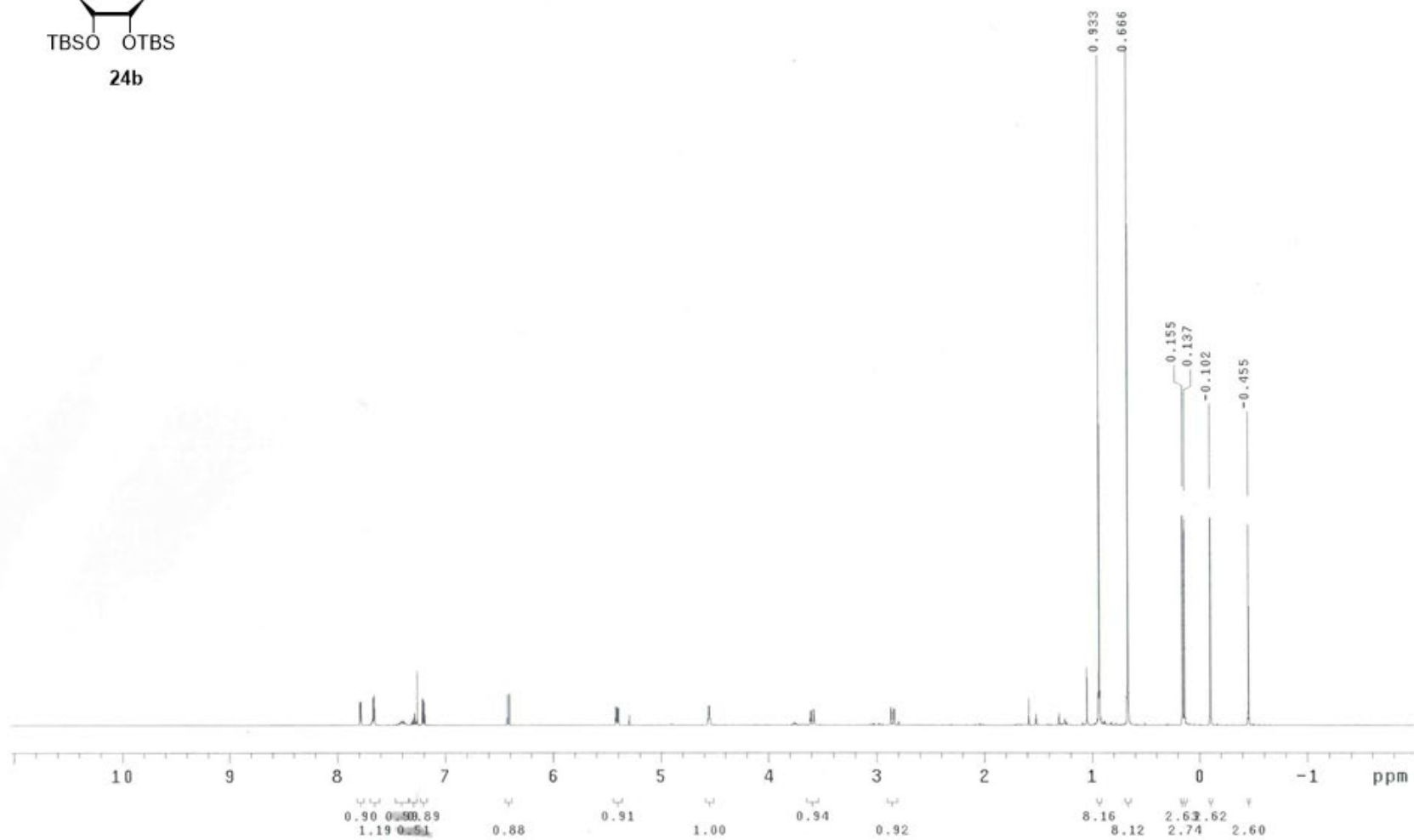
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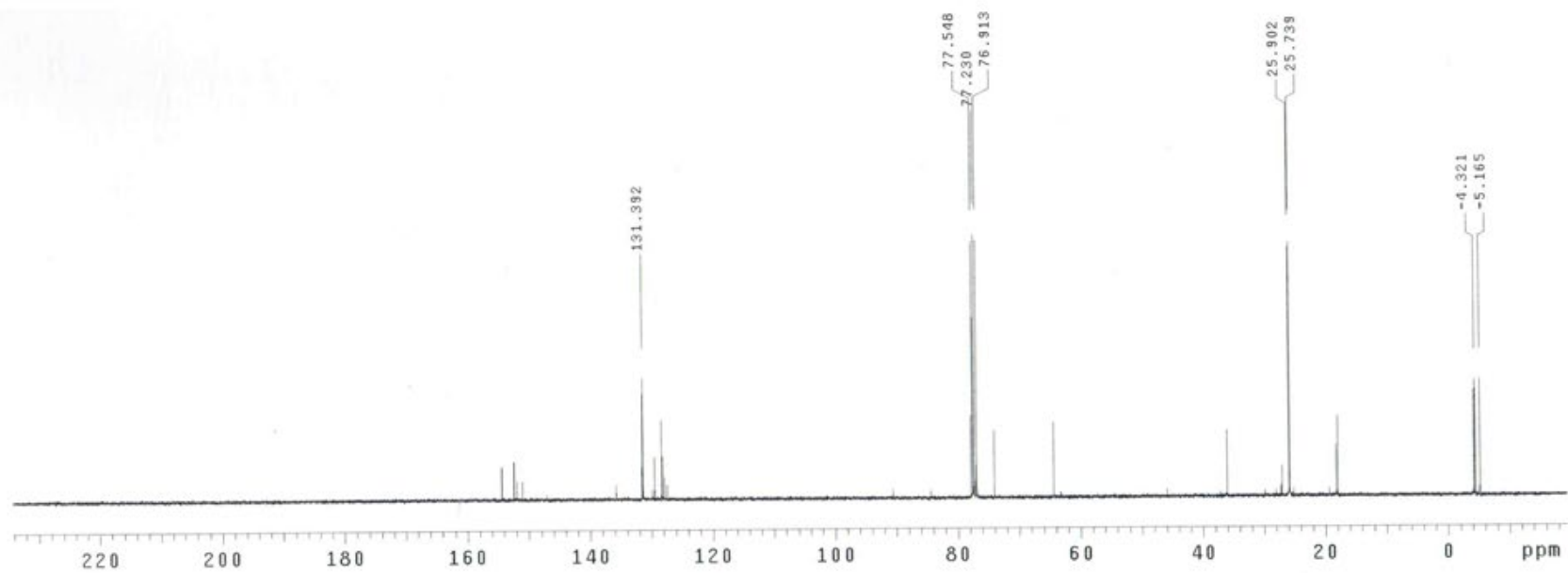
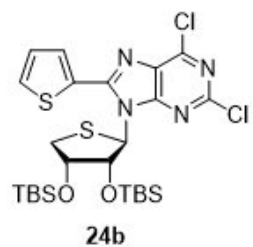


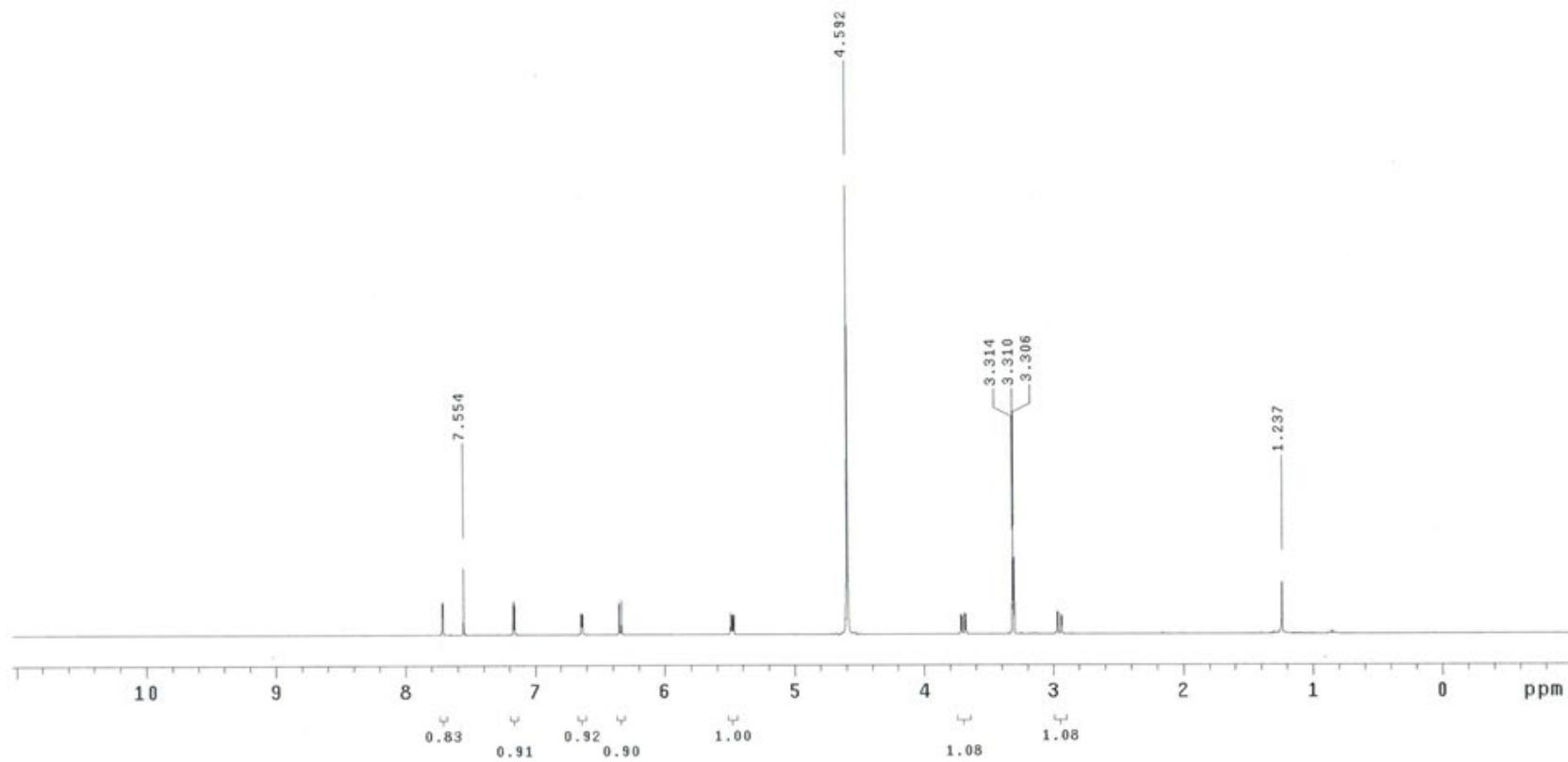
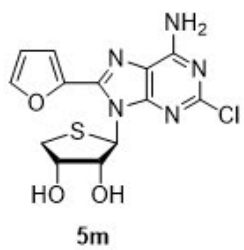


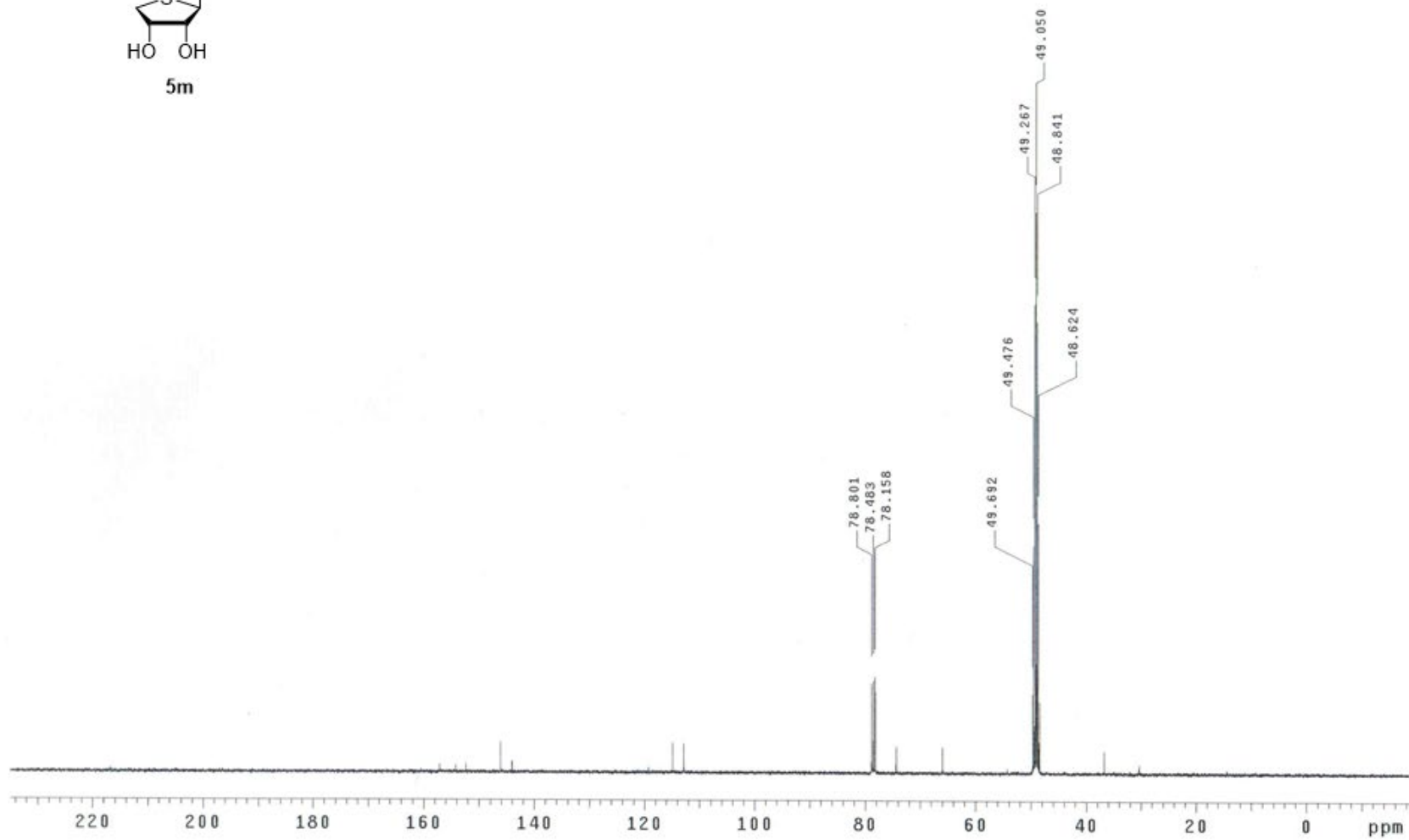
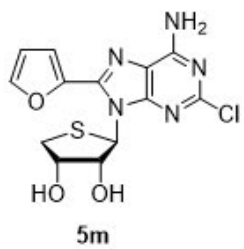
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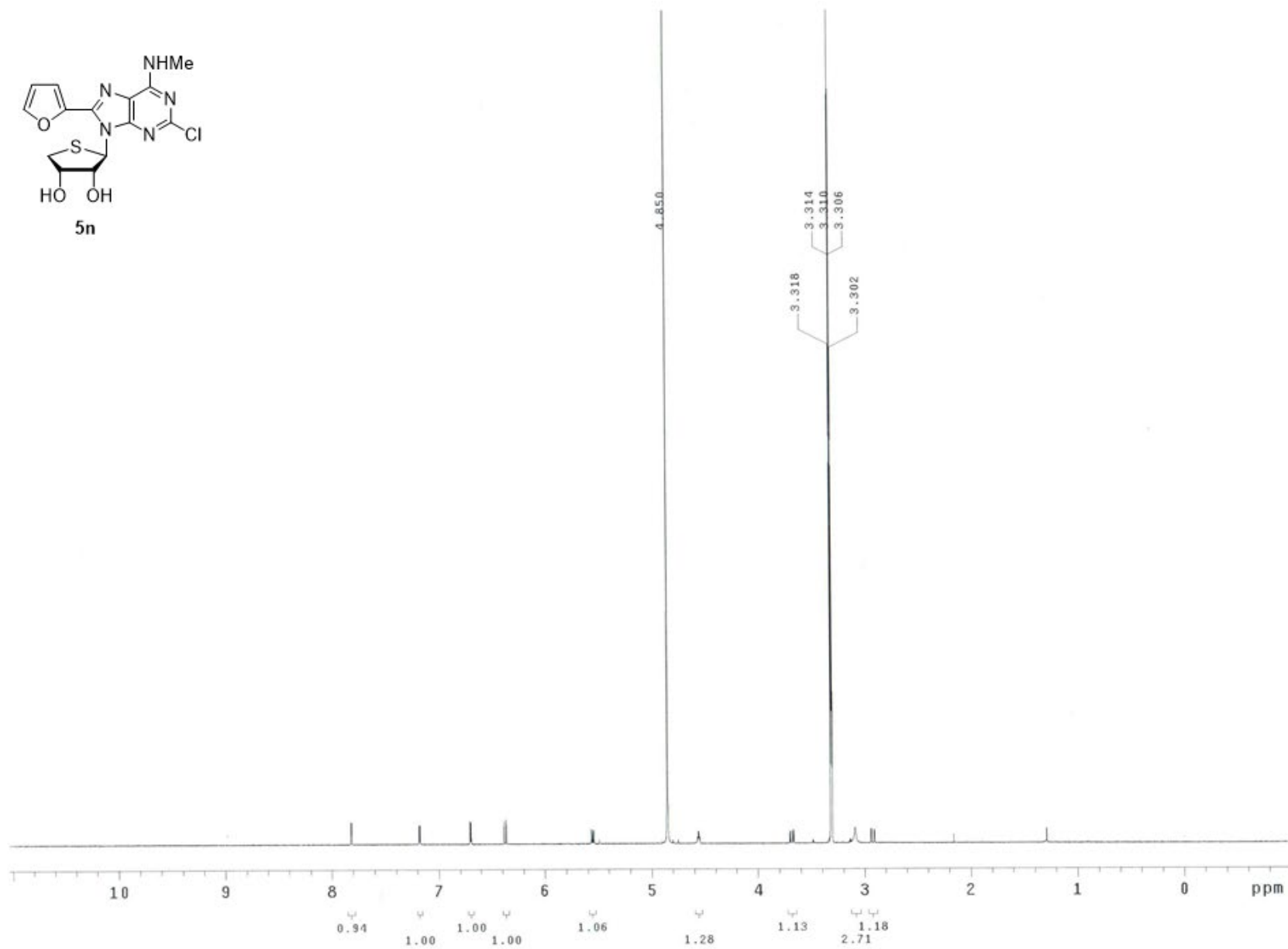
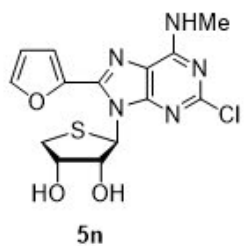


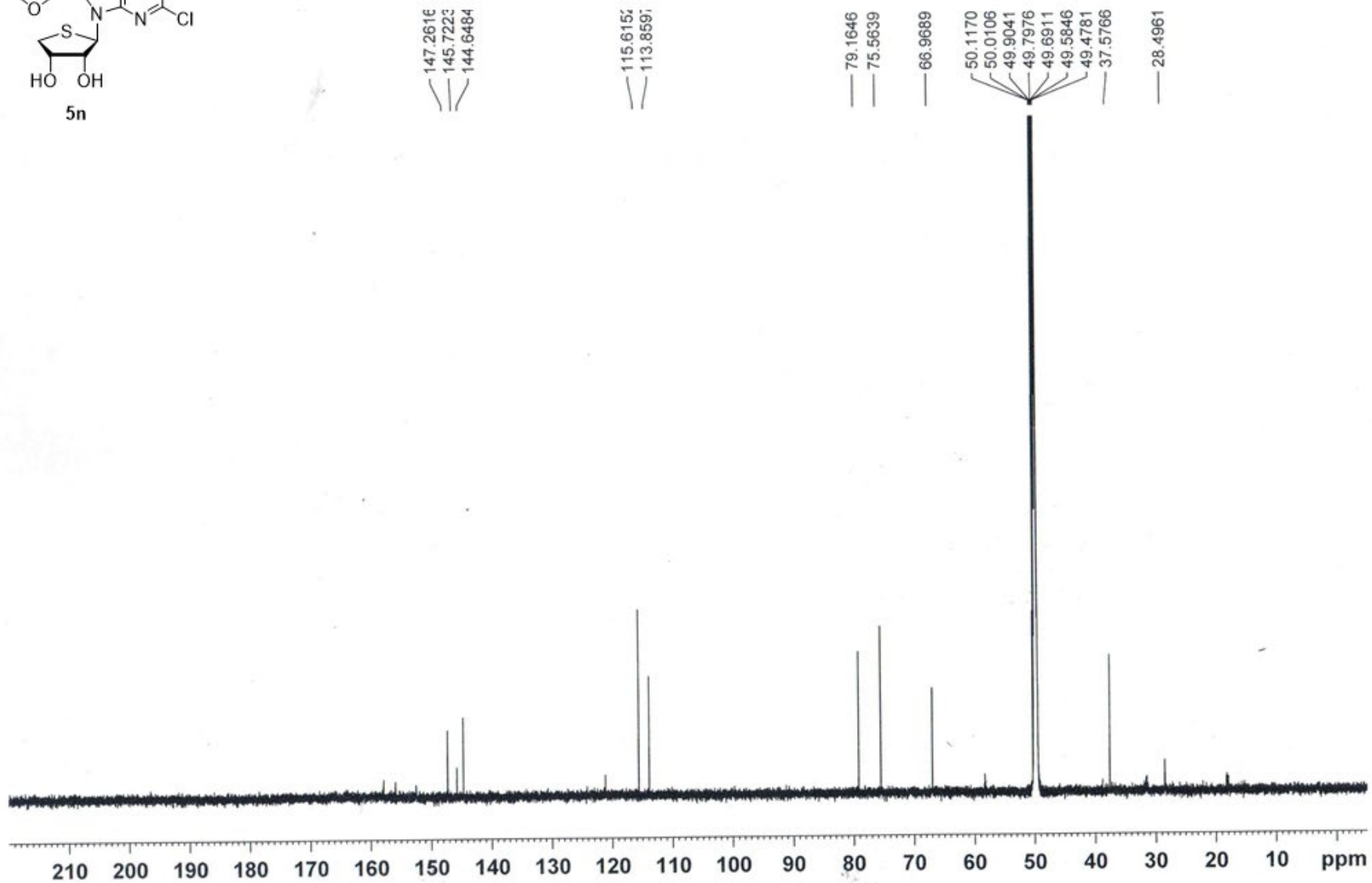
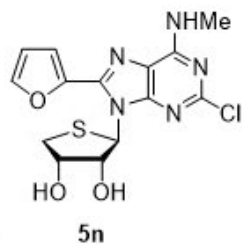


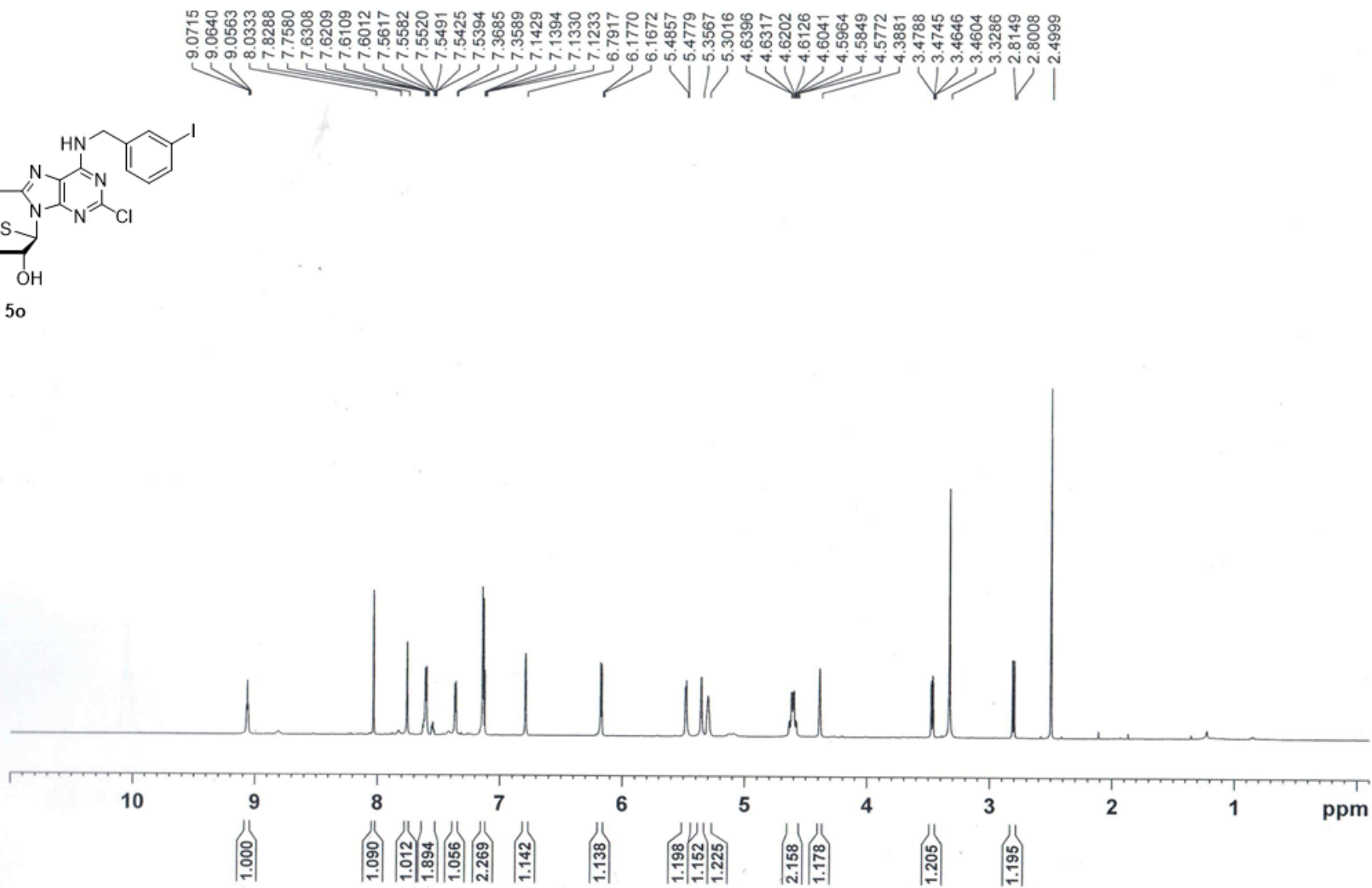
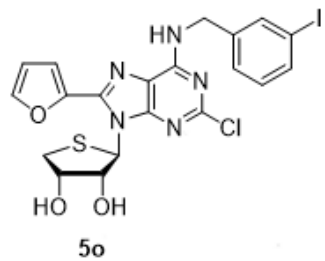


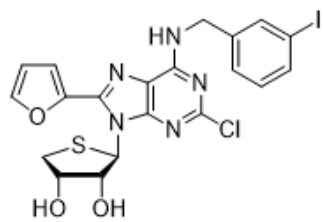




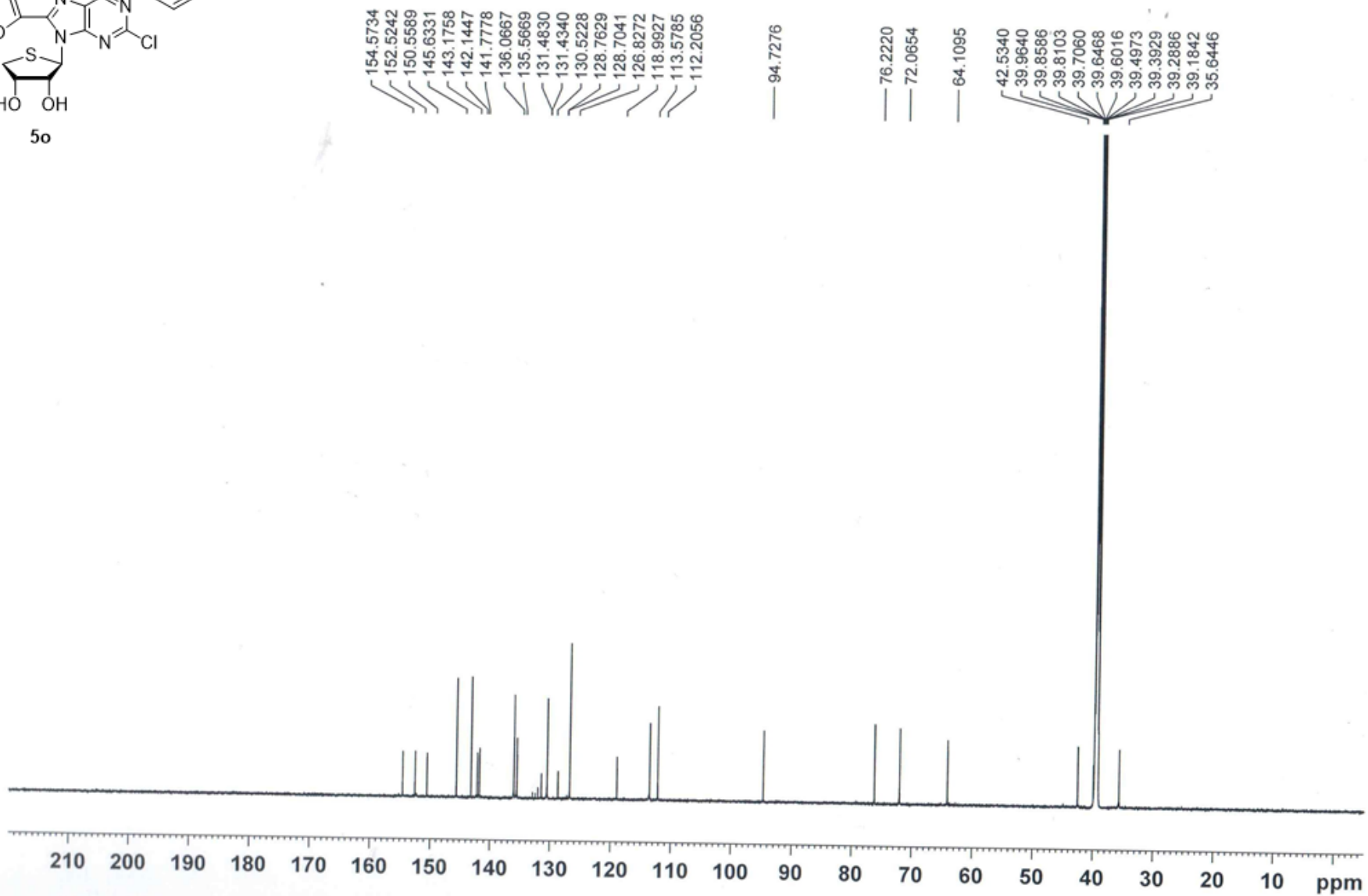


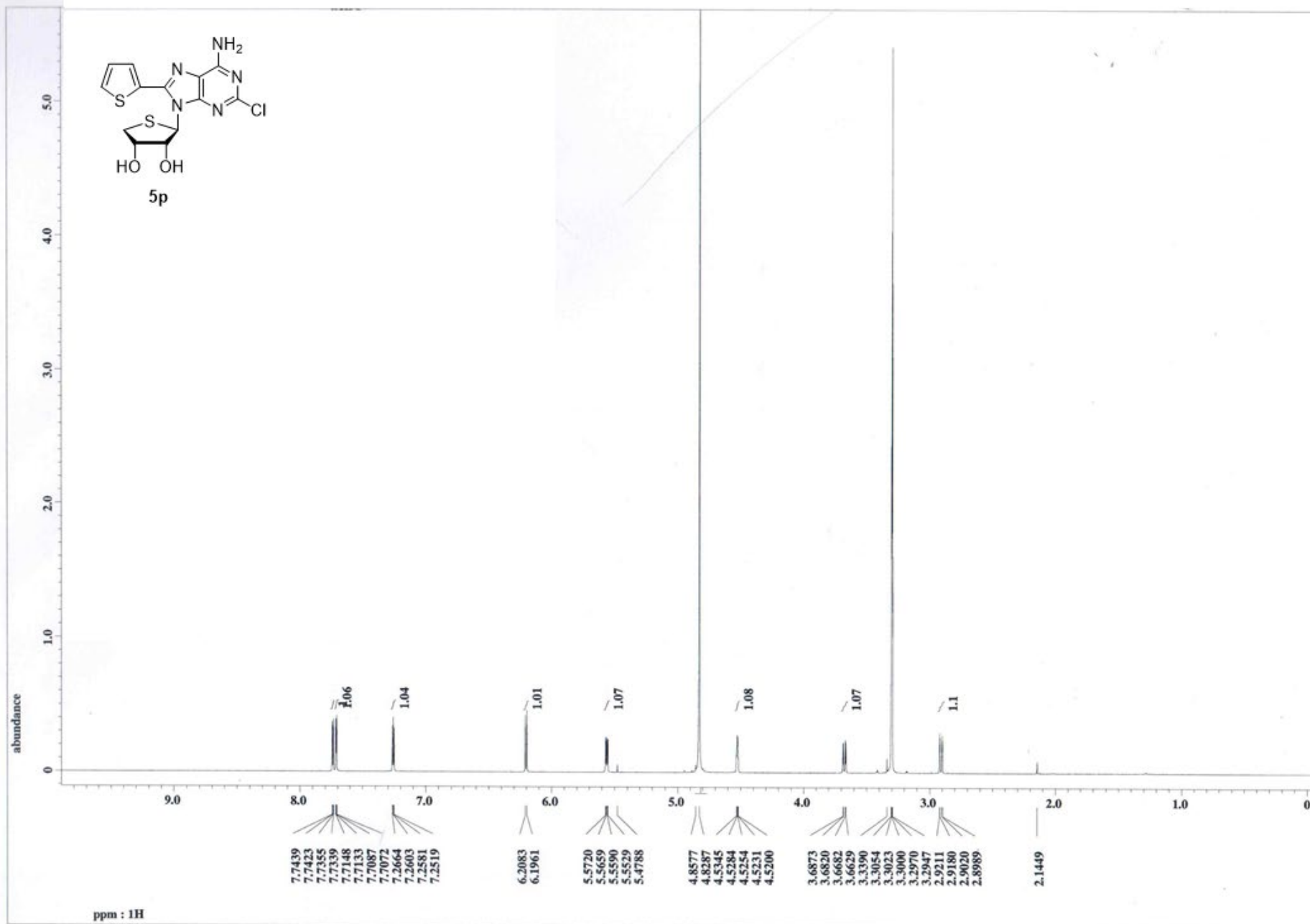


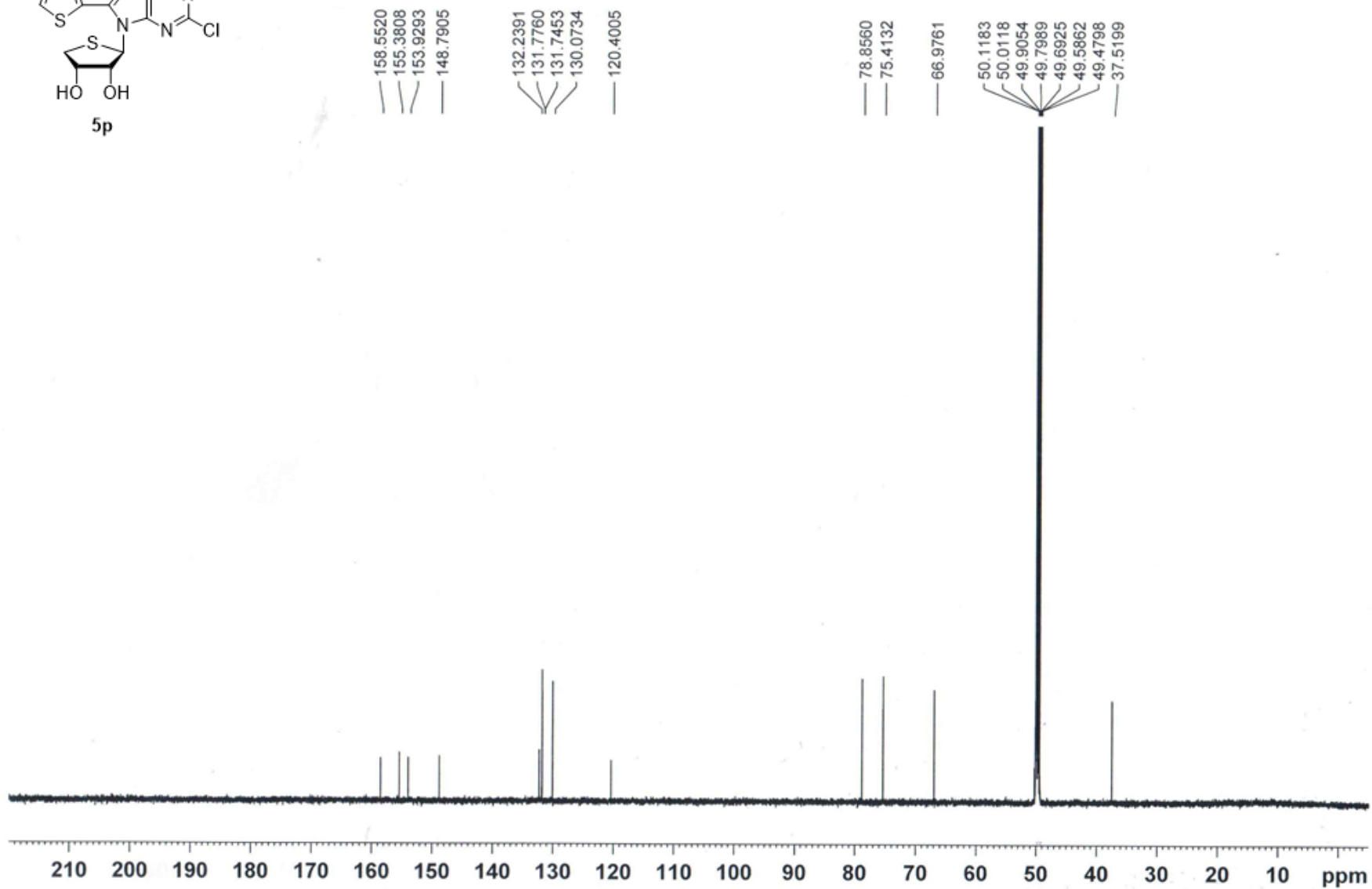
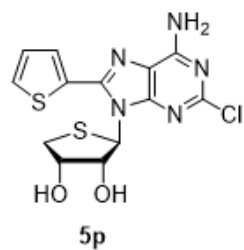


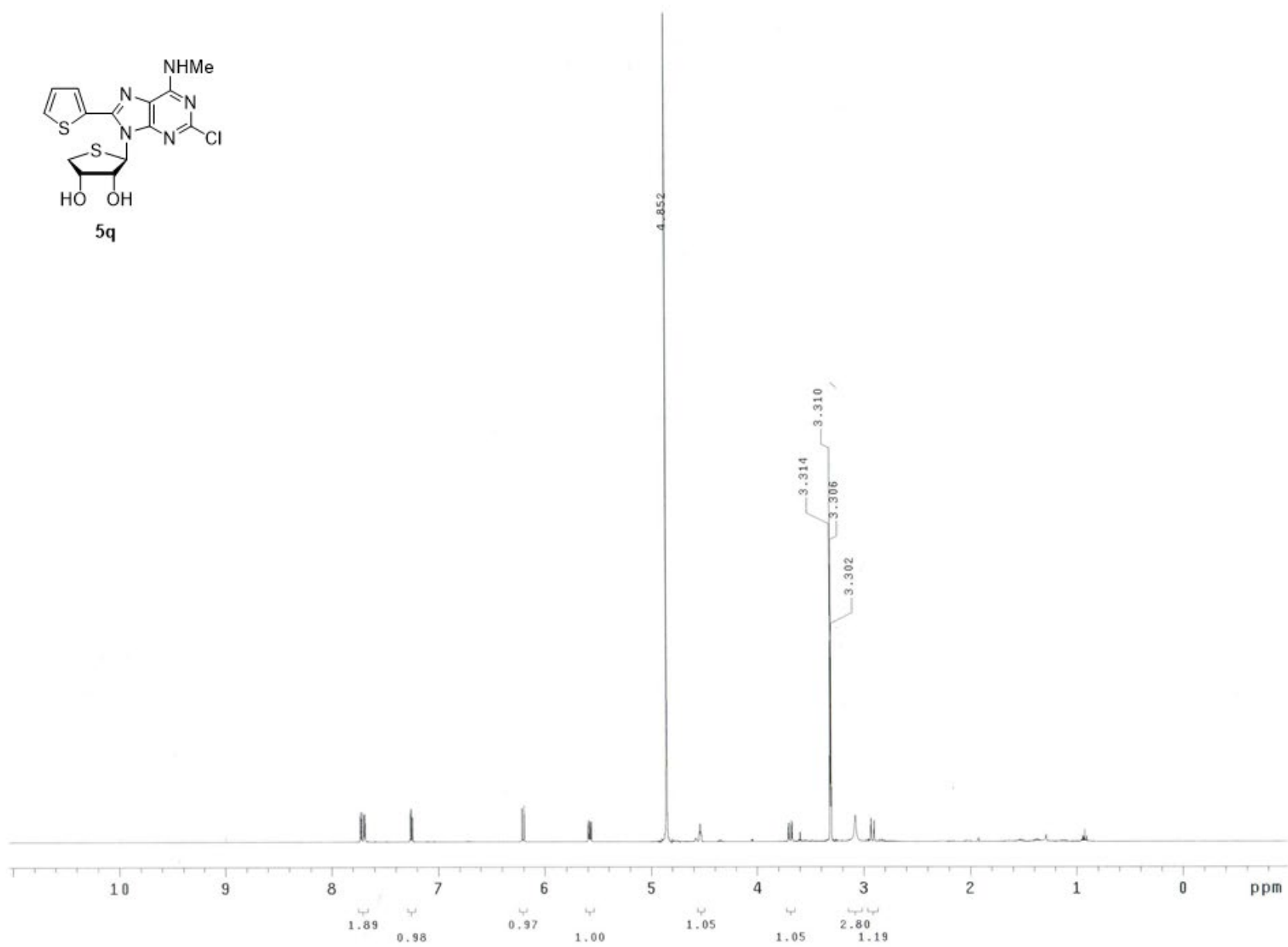
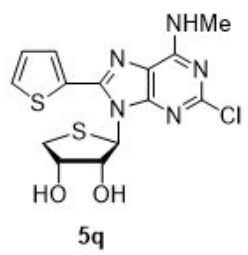


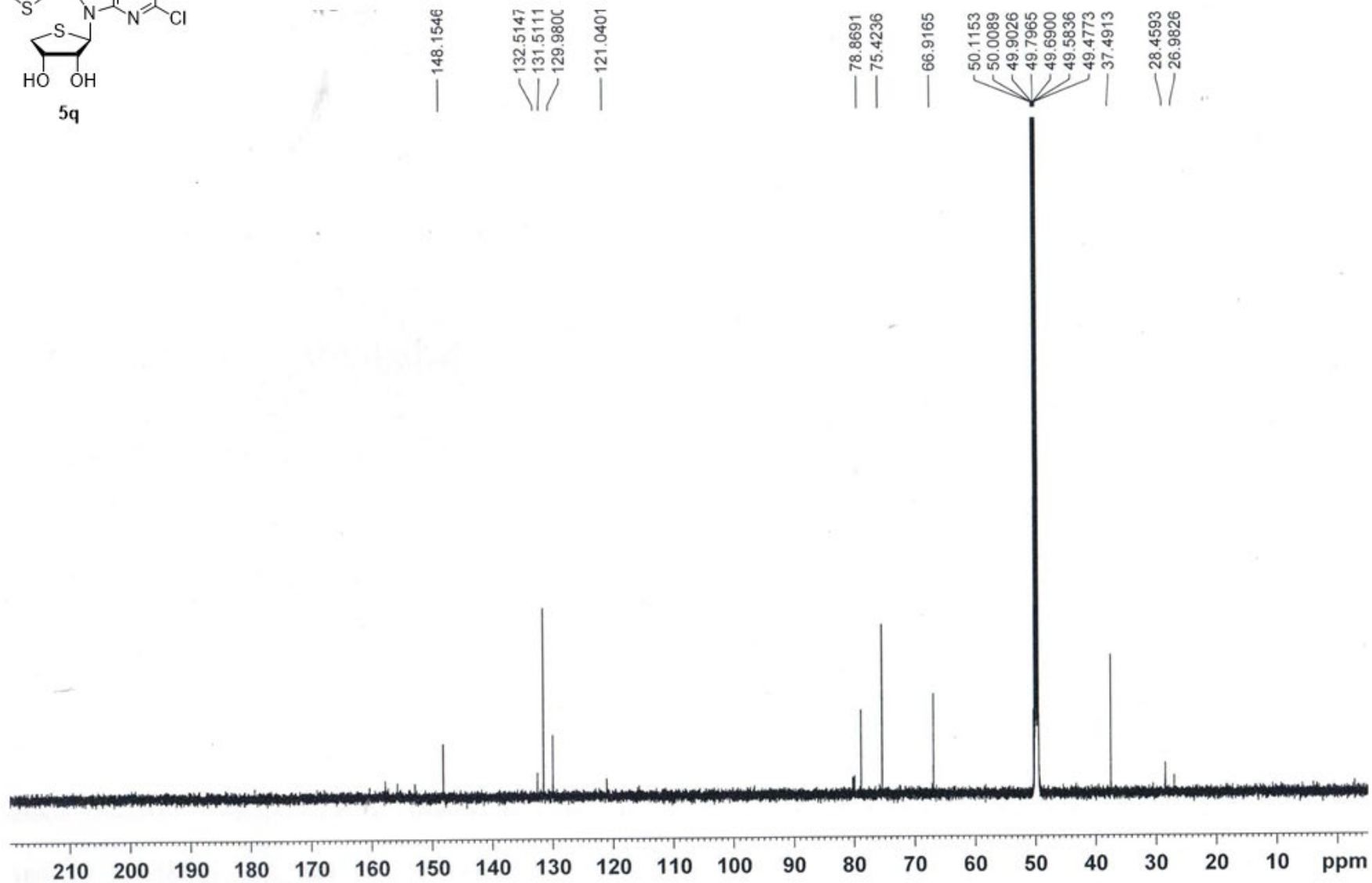
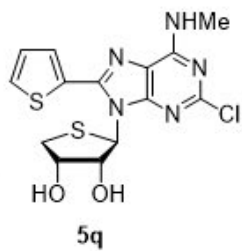
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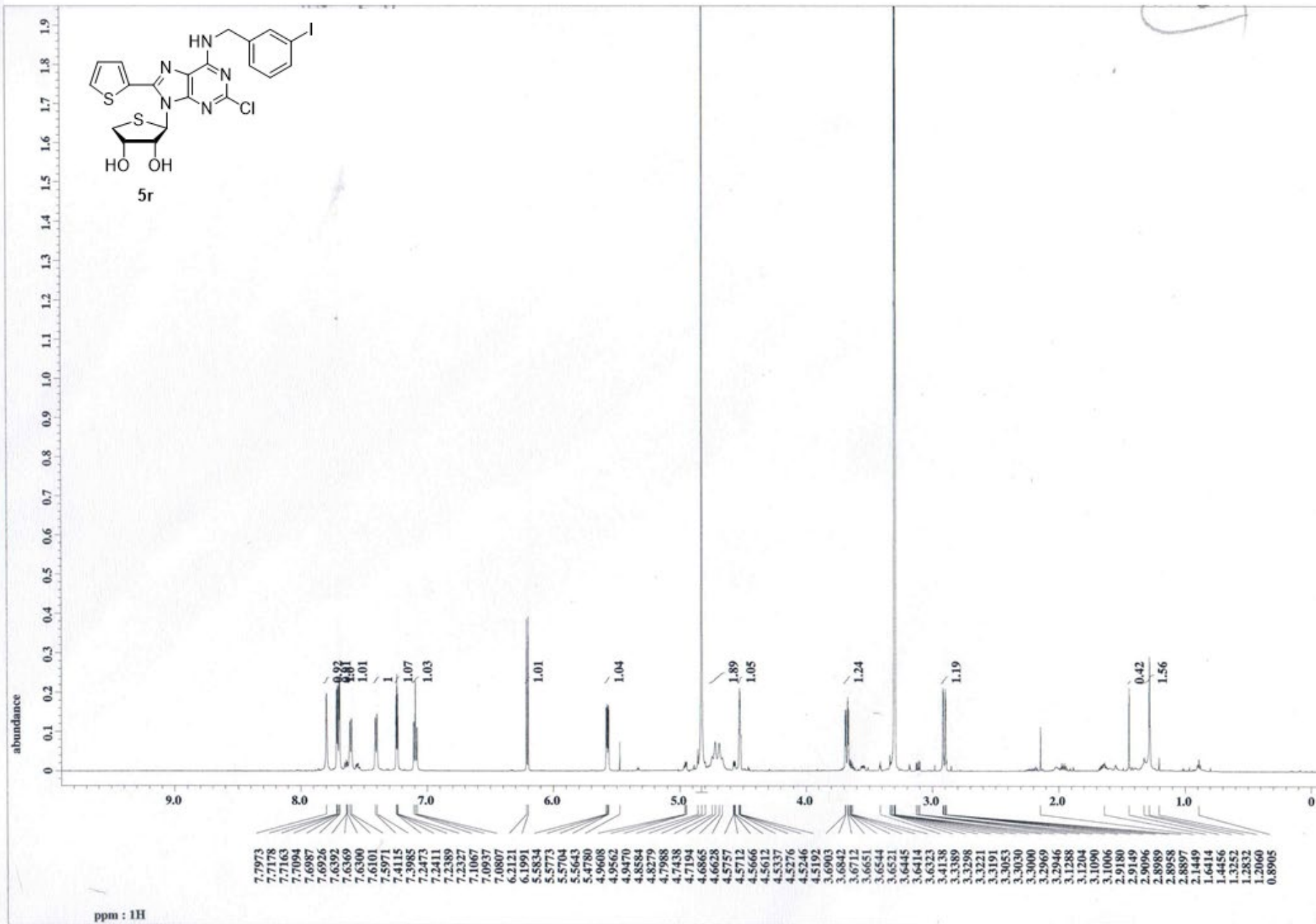


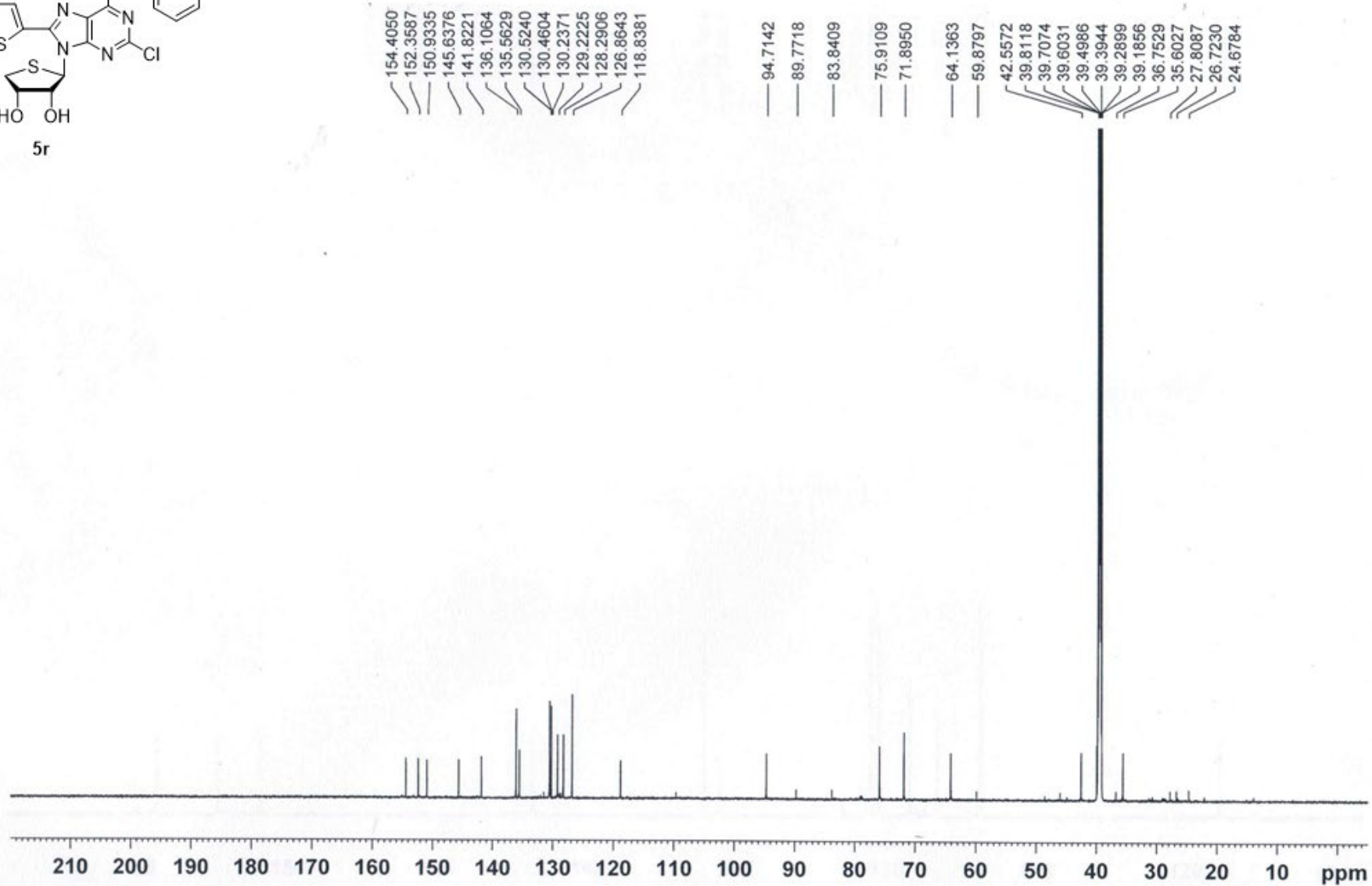
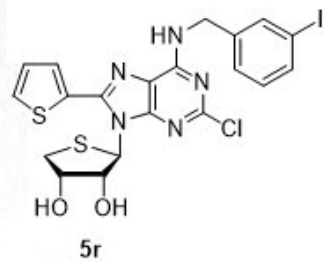


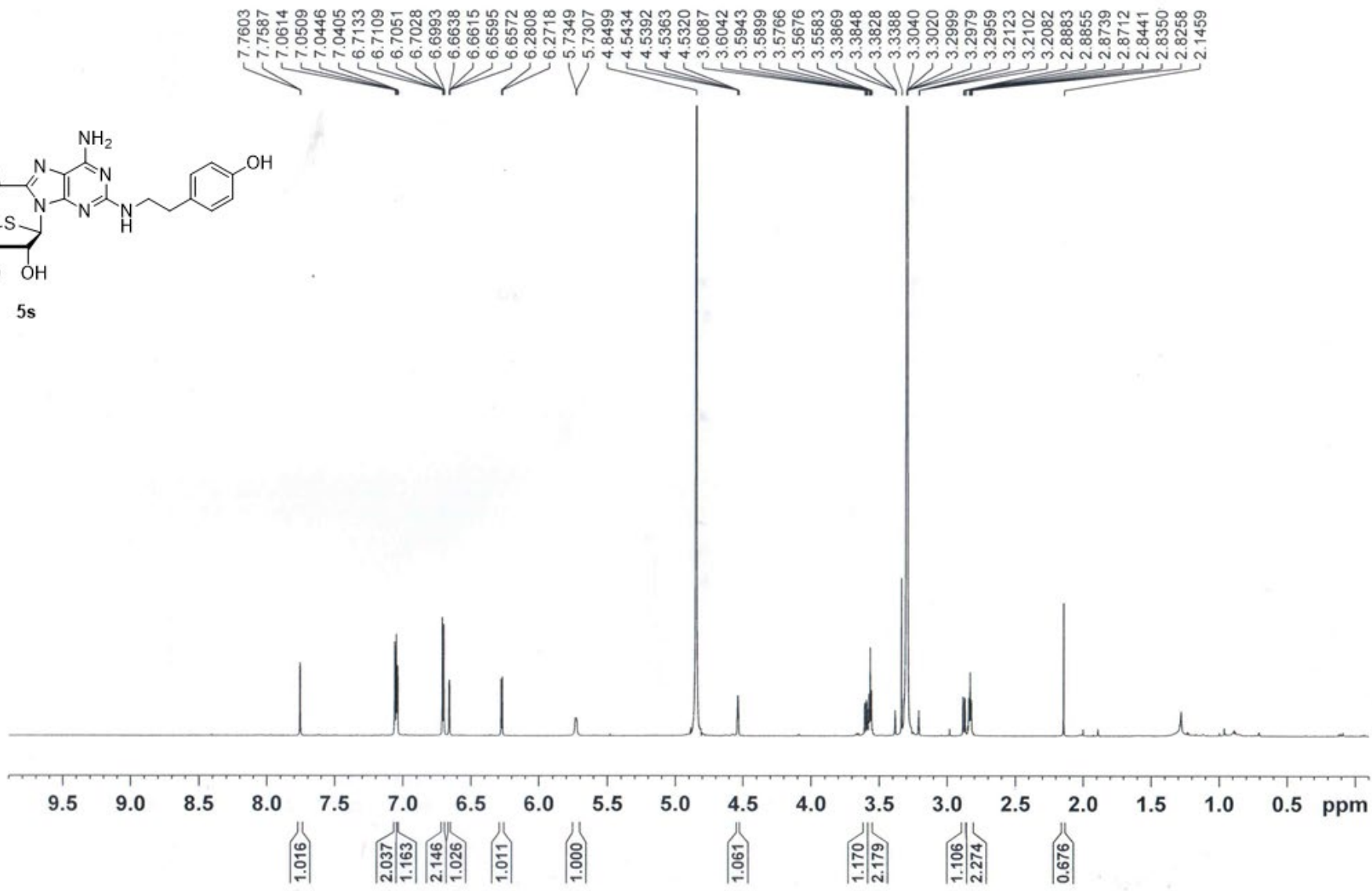
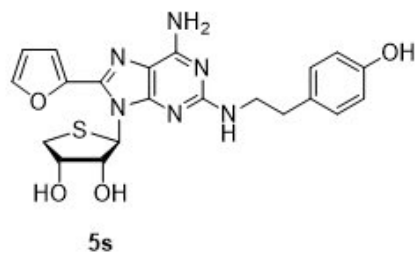


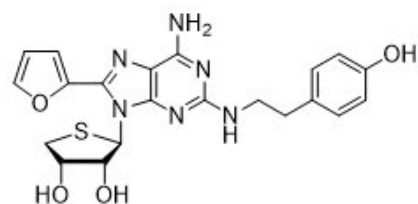












5s

158.7779
156.0930
155.4608

144.3447
144.2984

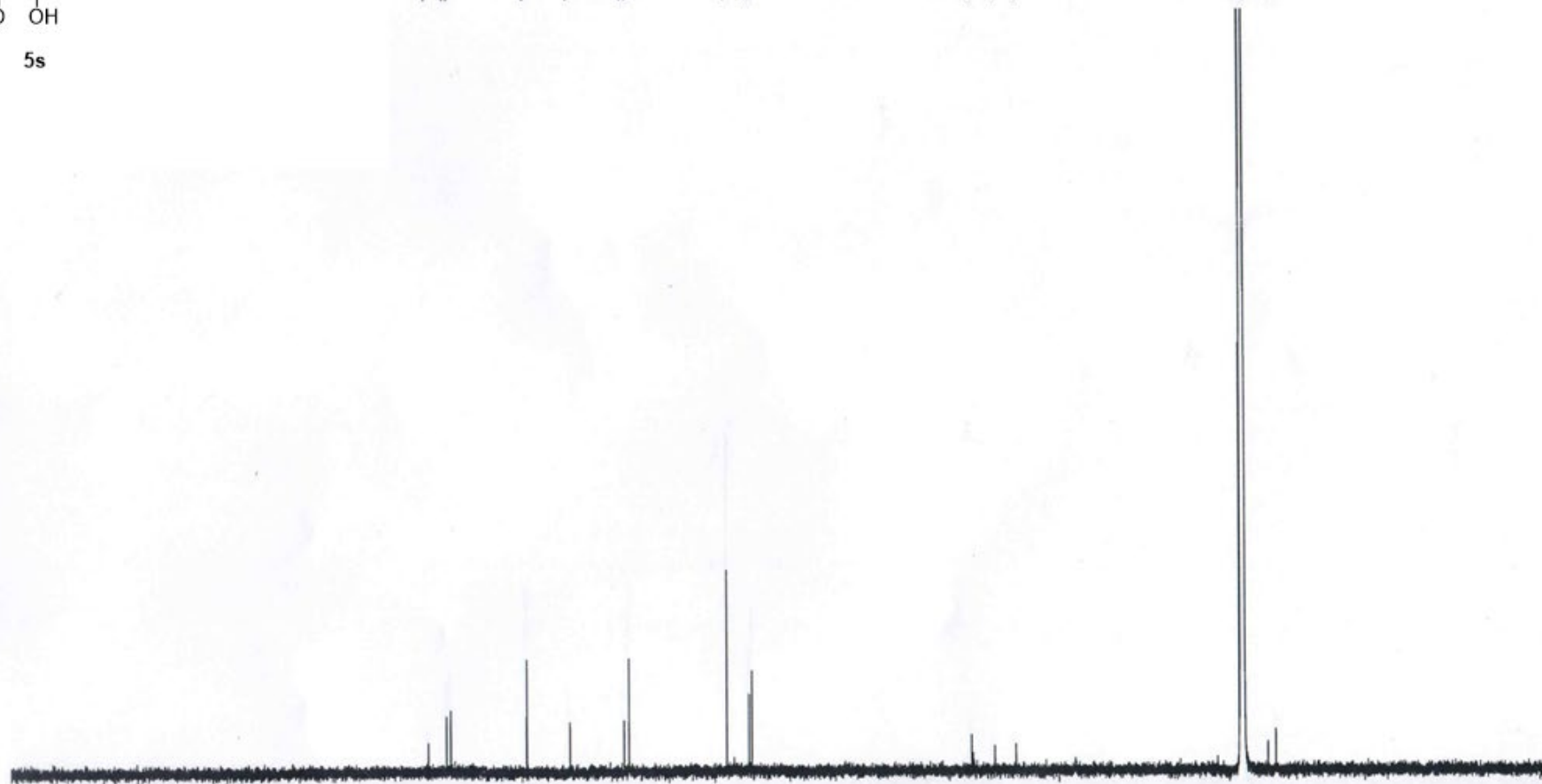
138.0123

130.0761
129.3702

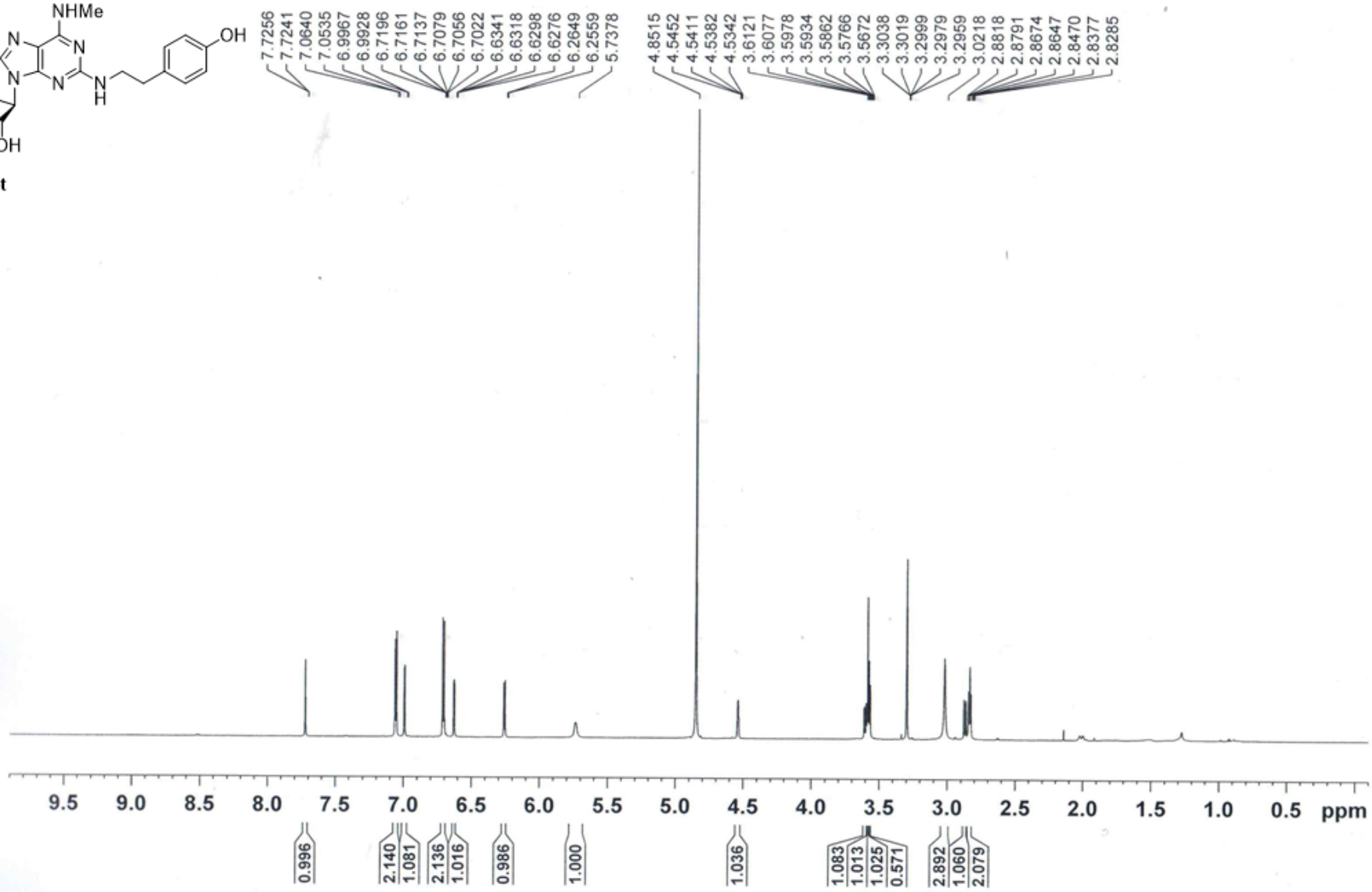
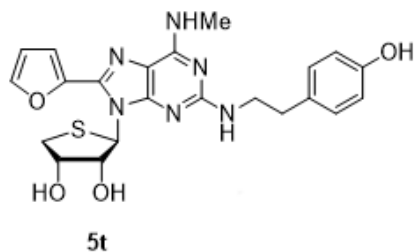
115.0541
111.7671
111.3414

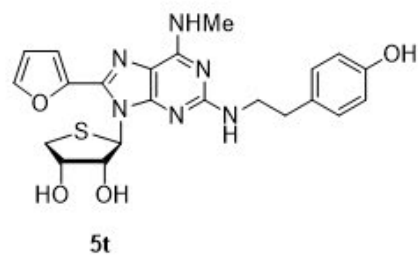
79.1527
78.9468
75.7526
72.6425

39.8123
39.7078
39.6036
39.4991
39.3948
39.2904
39.1860
35.7532
34.5866

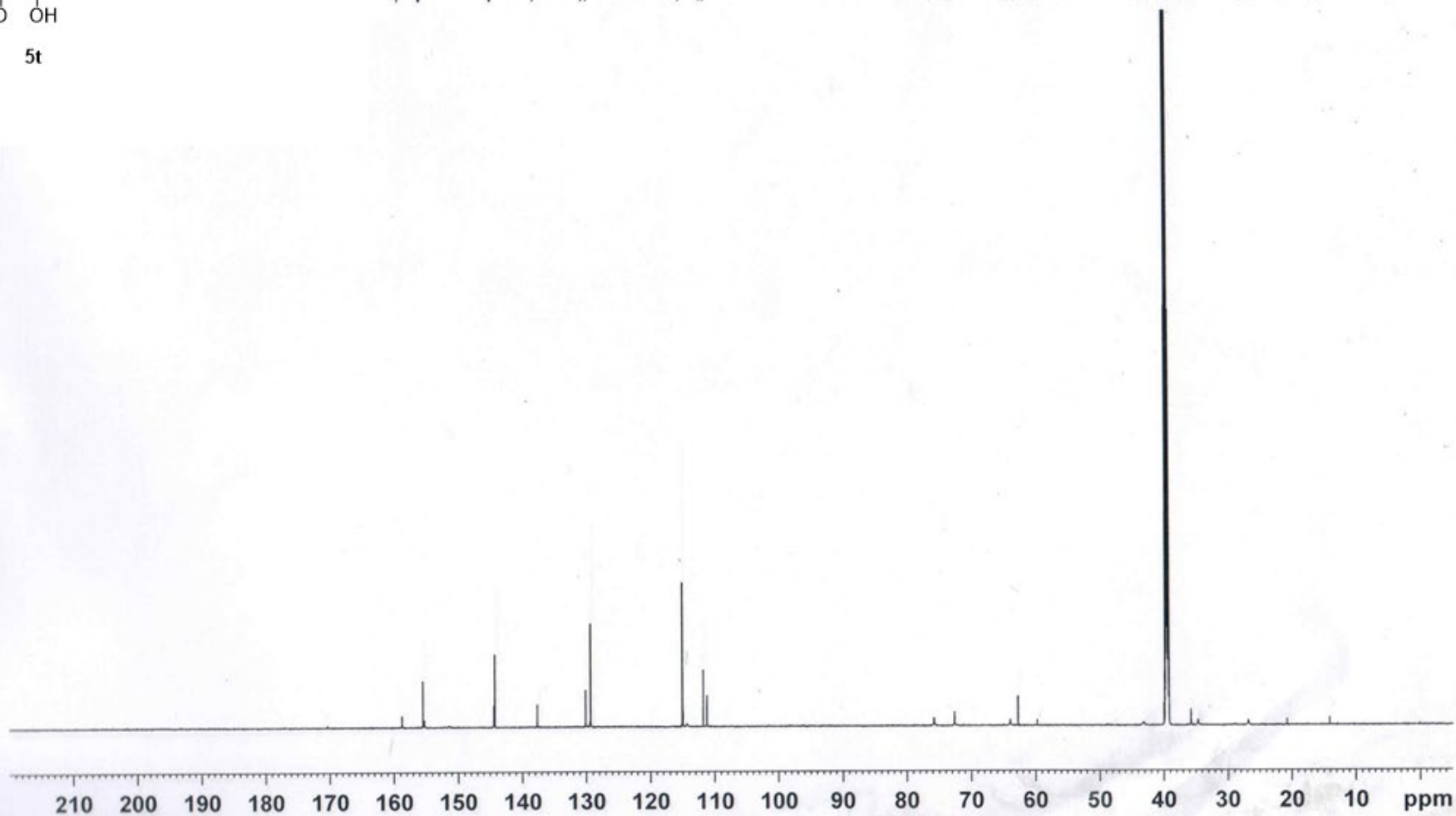


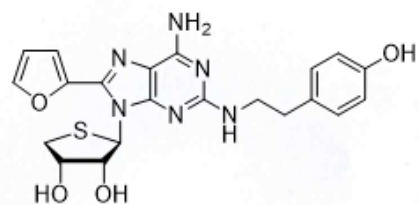
210 200 190 180 170 160 150 140 130 120 110 100 90 80 70 60 50 40 30 20 10 ppm



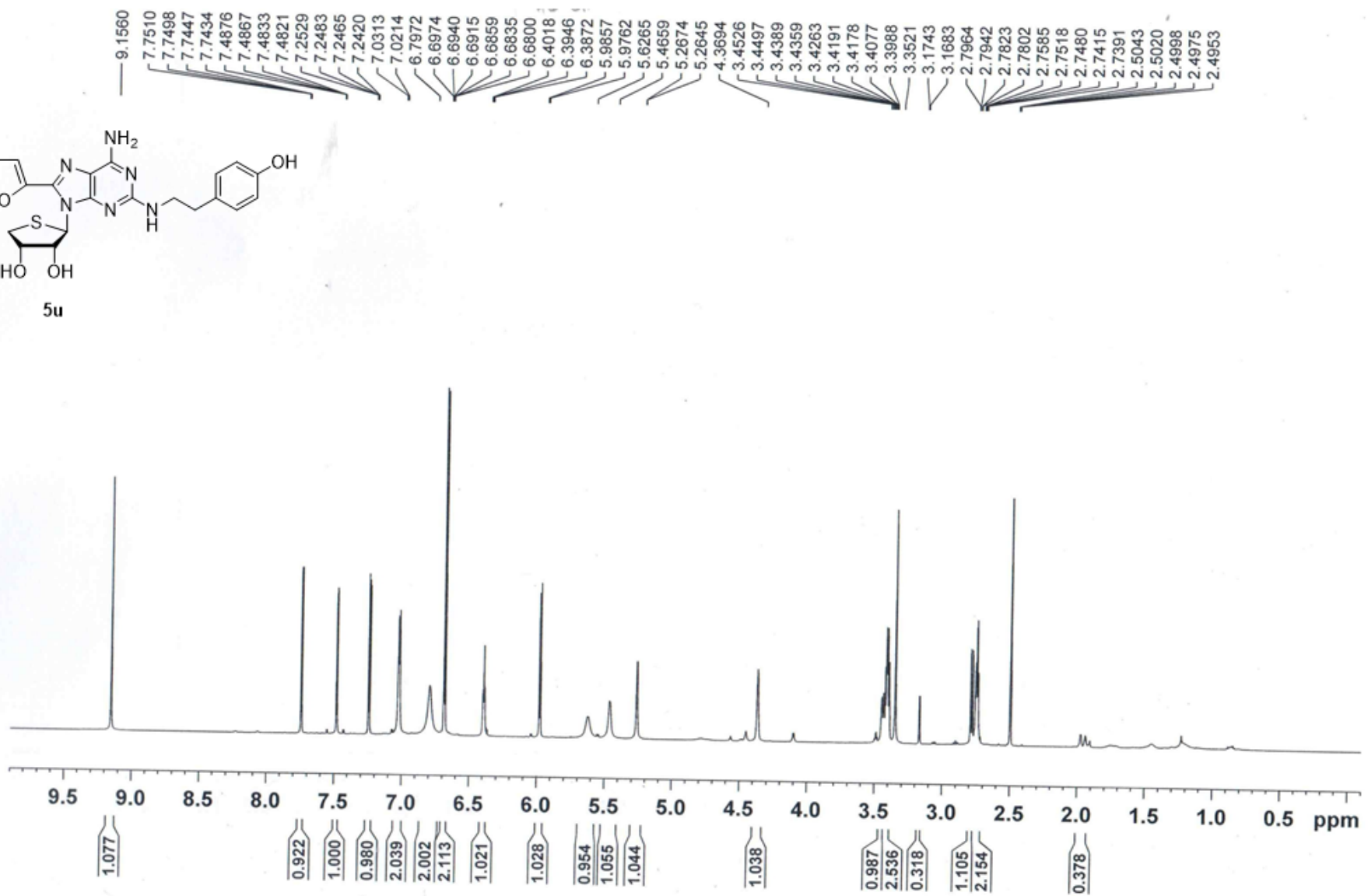


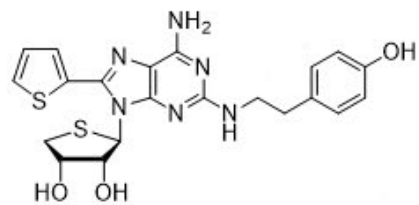
- 158.7593
- 155.4645
- 155.3246
- 155.2552
- 144.4007
- 144.2425
- 137.6672
- 130.0897
- 129.3446
- 115.0838
- 111.7744
- 111.1909
- 75.8287
- 72.6451
- 63.9635
- 62.7699
- 59.7349
- 43.1250
- 39.8094
- 39.7050
- 39.6007
- 39.4963
- 39.3919
- 39.2876
- 39.1831
- 35.7641
- 34.6425
- 26.8084
- 20.7456
- 14.0667



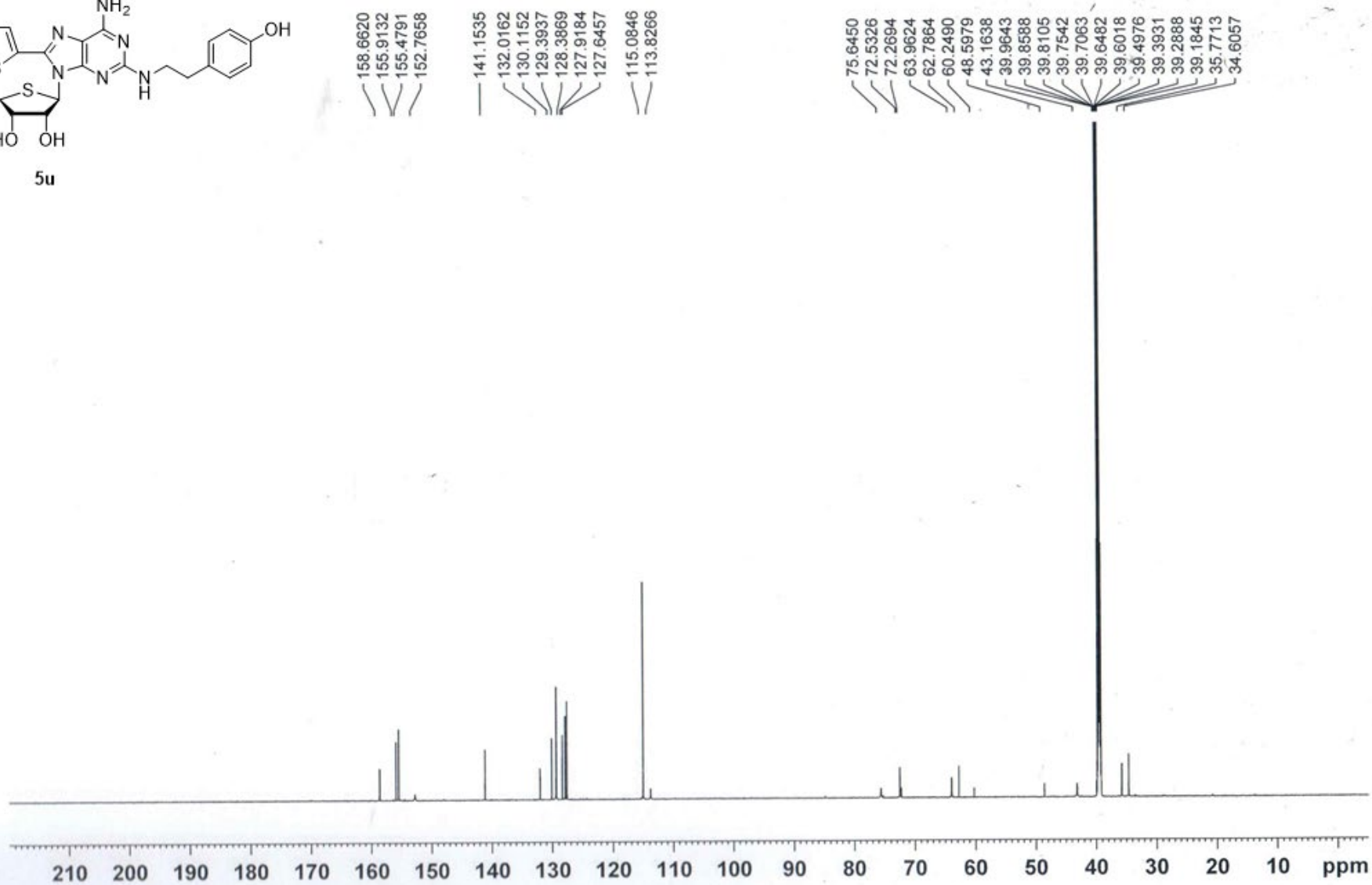


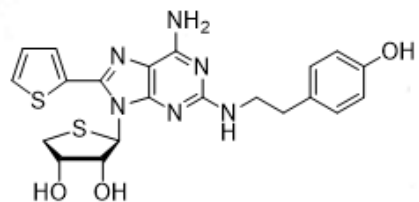
5u





5u





5u

