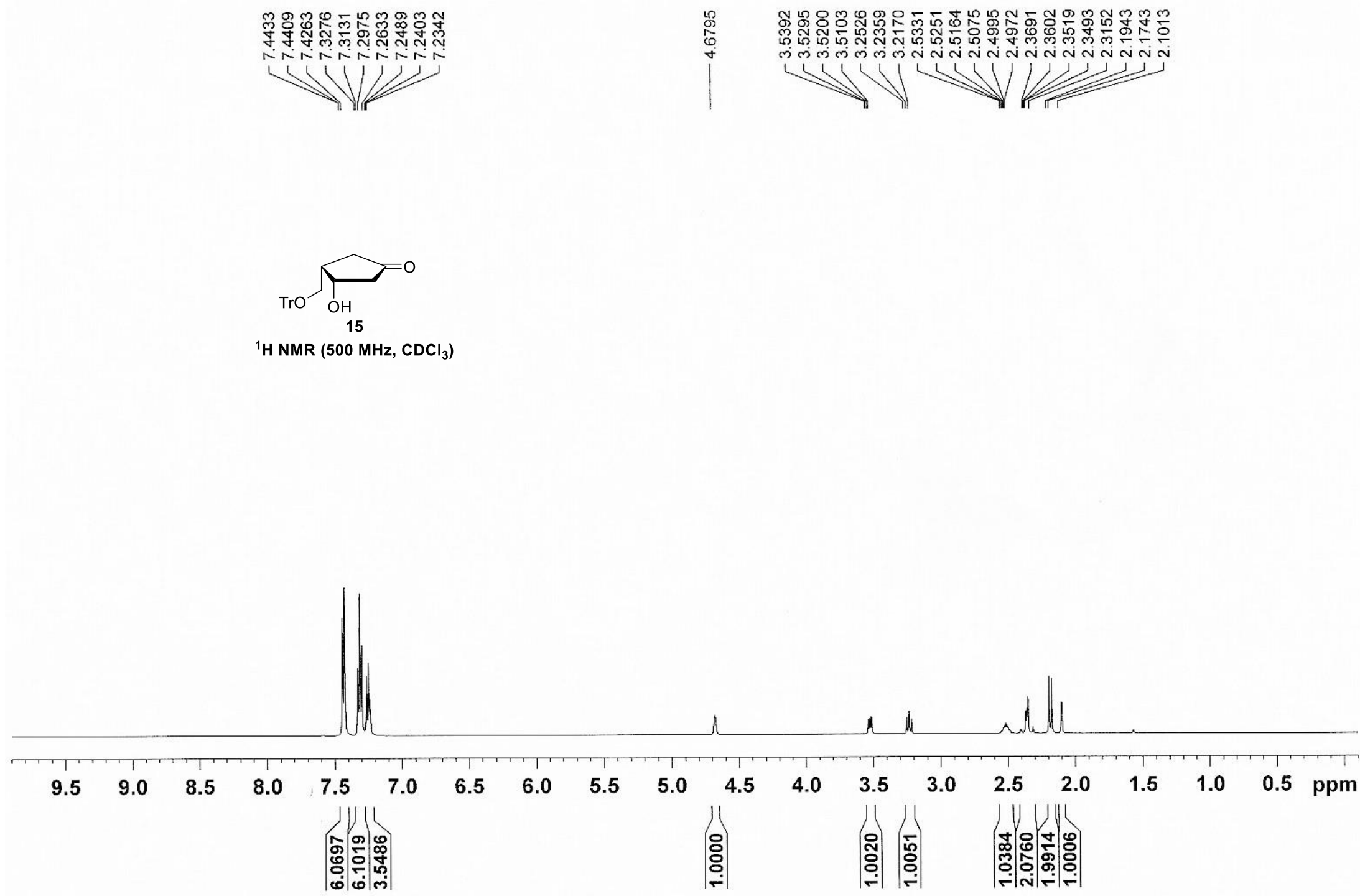
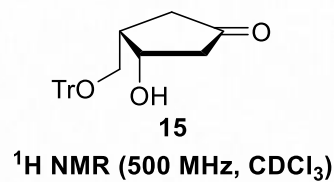


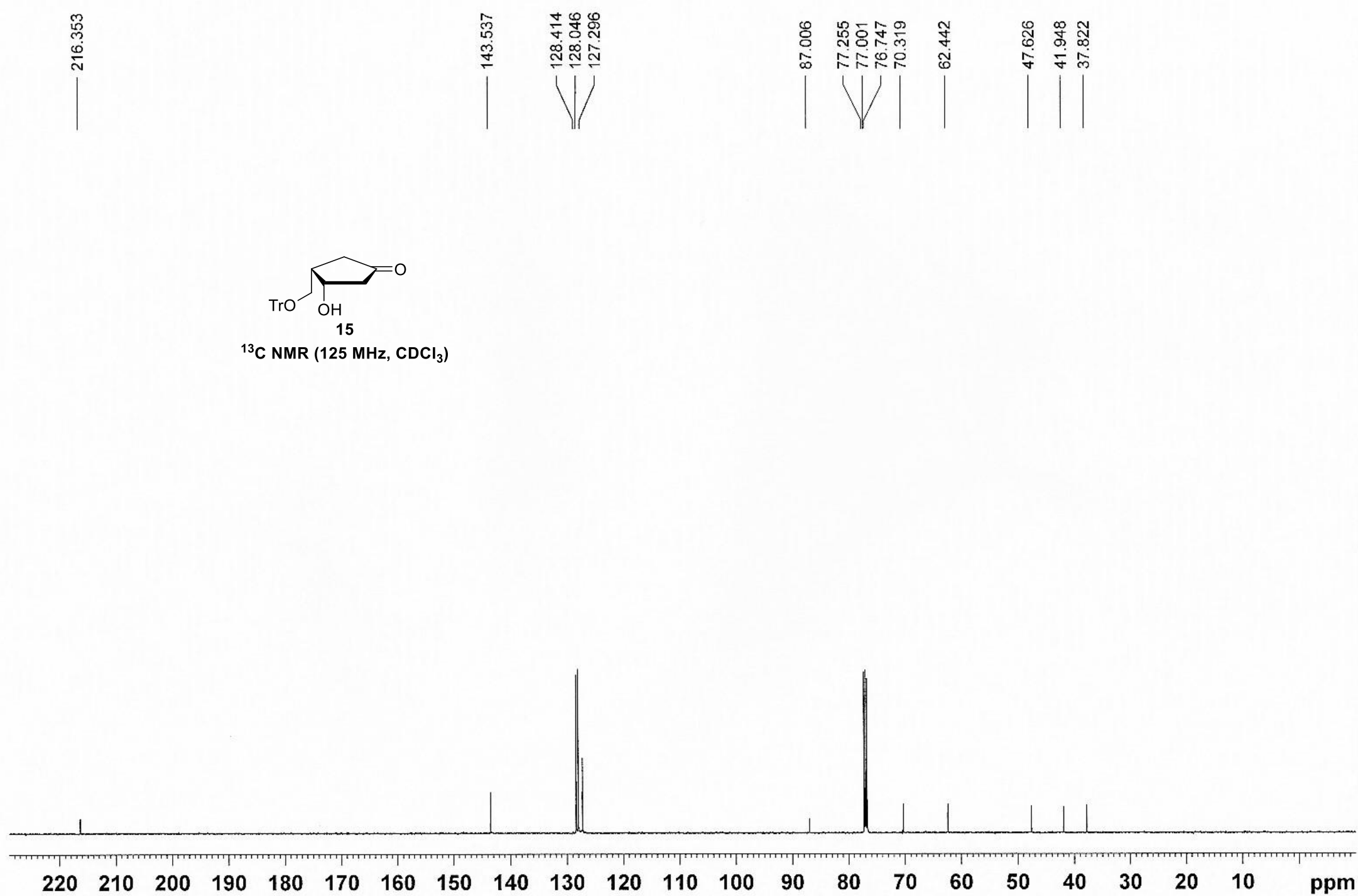
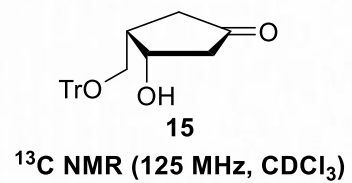
Supporting Information

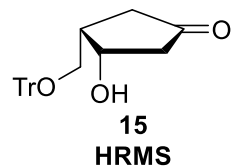
An Alternative and Efficient Synthesis of MLN4924, A Selective NEDD8 Inhibitor

Hong-Rae Kim,^a Young Eum Hyun,^a Dnyandev B. Jarhad,^a Jinha Yu,^a and Lak Shin Jeong^{*a}

^aResearch Institute of Pharmaceutical Sciences, College of Pharmacy, Seoul National University,
Seoul 08826, Korea, E-mail: lakjeong@snu.ac.kr; Tel: +82-2-880-7850.







Data : T0404003 Date : 04-Apr-2019 08:42

Instrument : MStation

Sample : A

Note : m-NBA

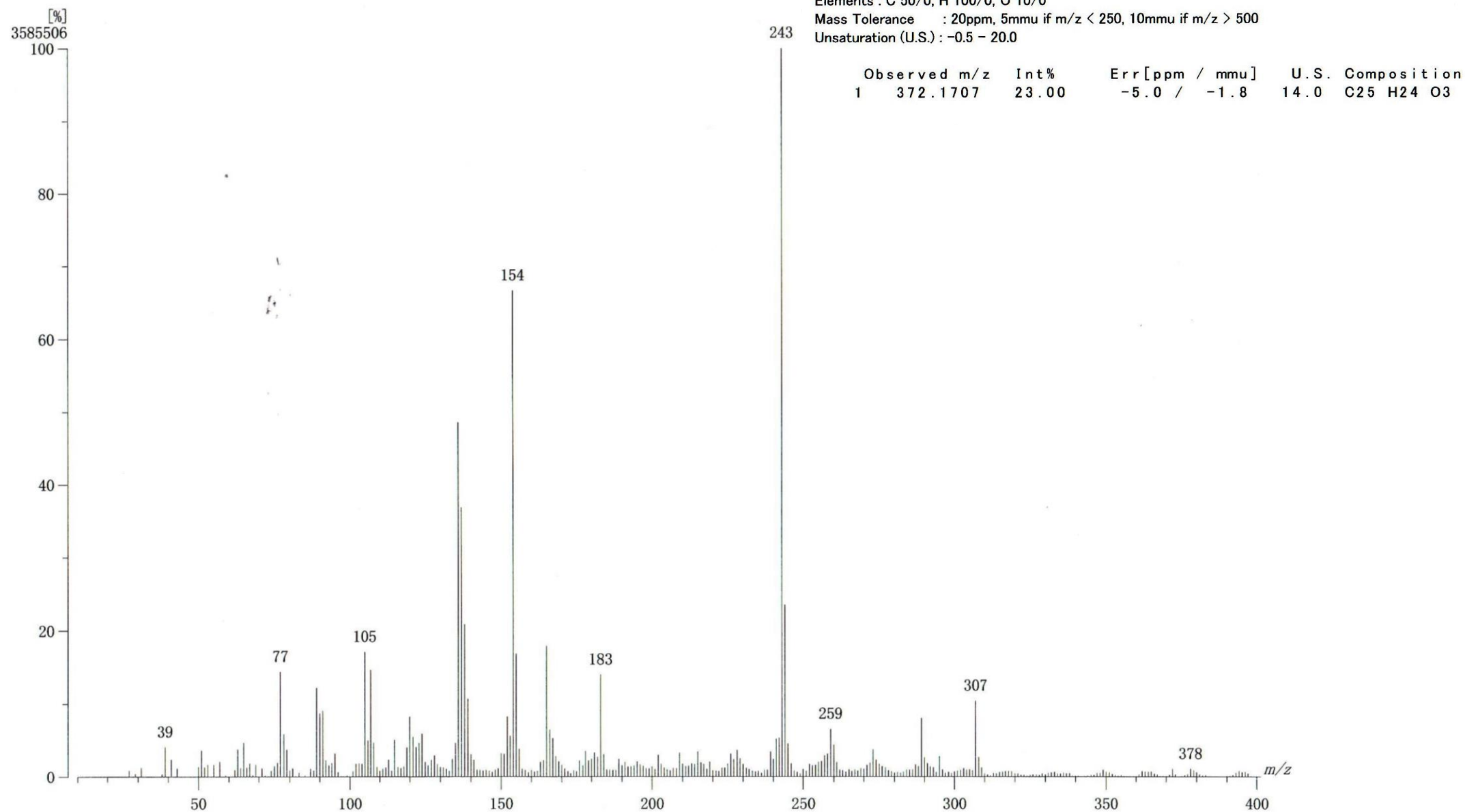
Inlet : Direct Ion Mode : FAB+

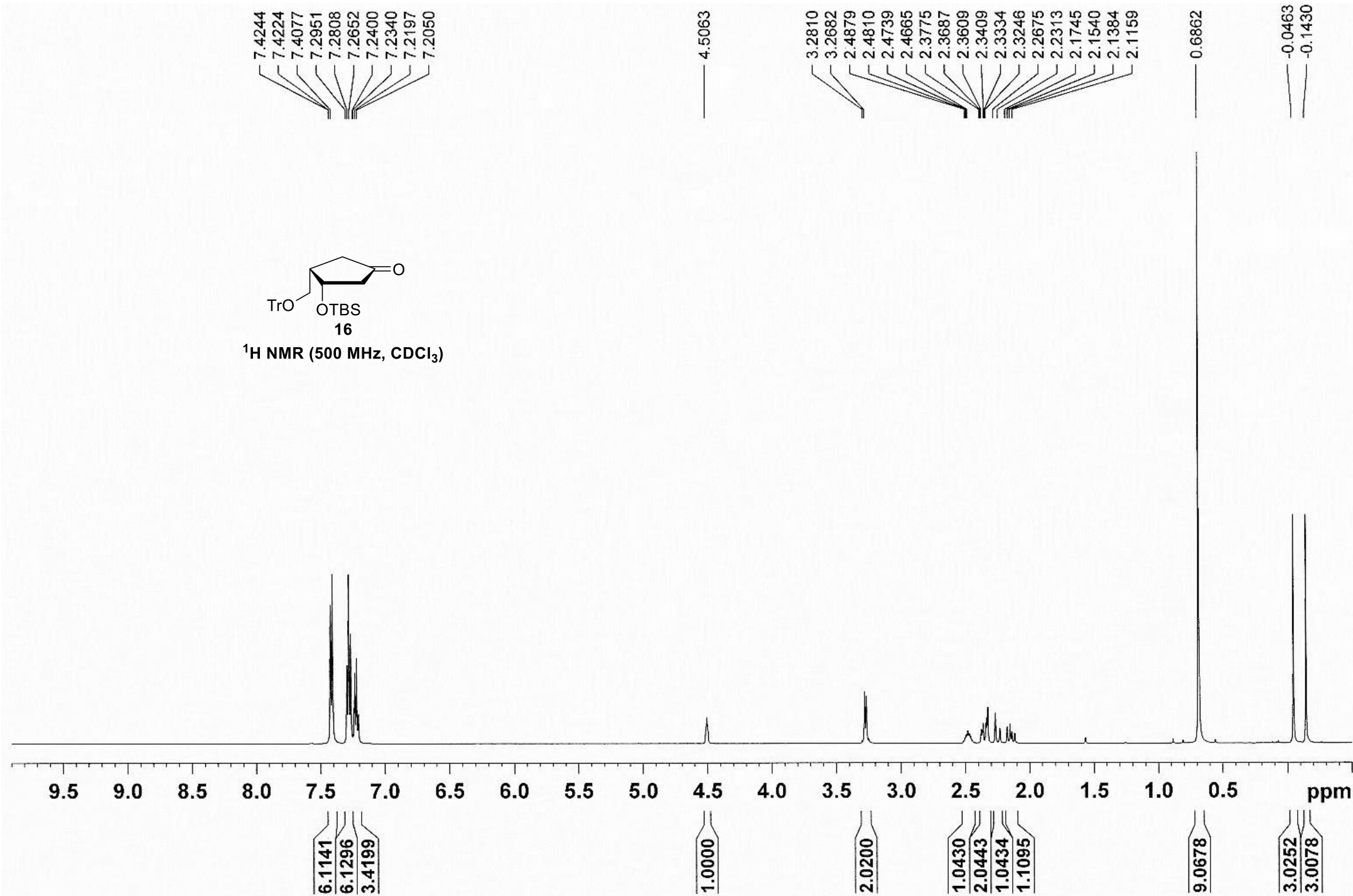
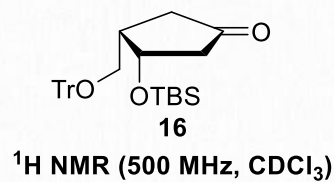
RT : 3.44 min Scan# : (60,66)+(16,21)

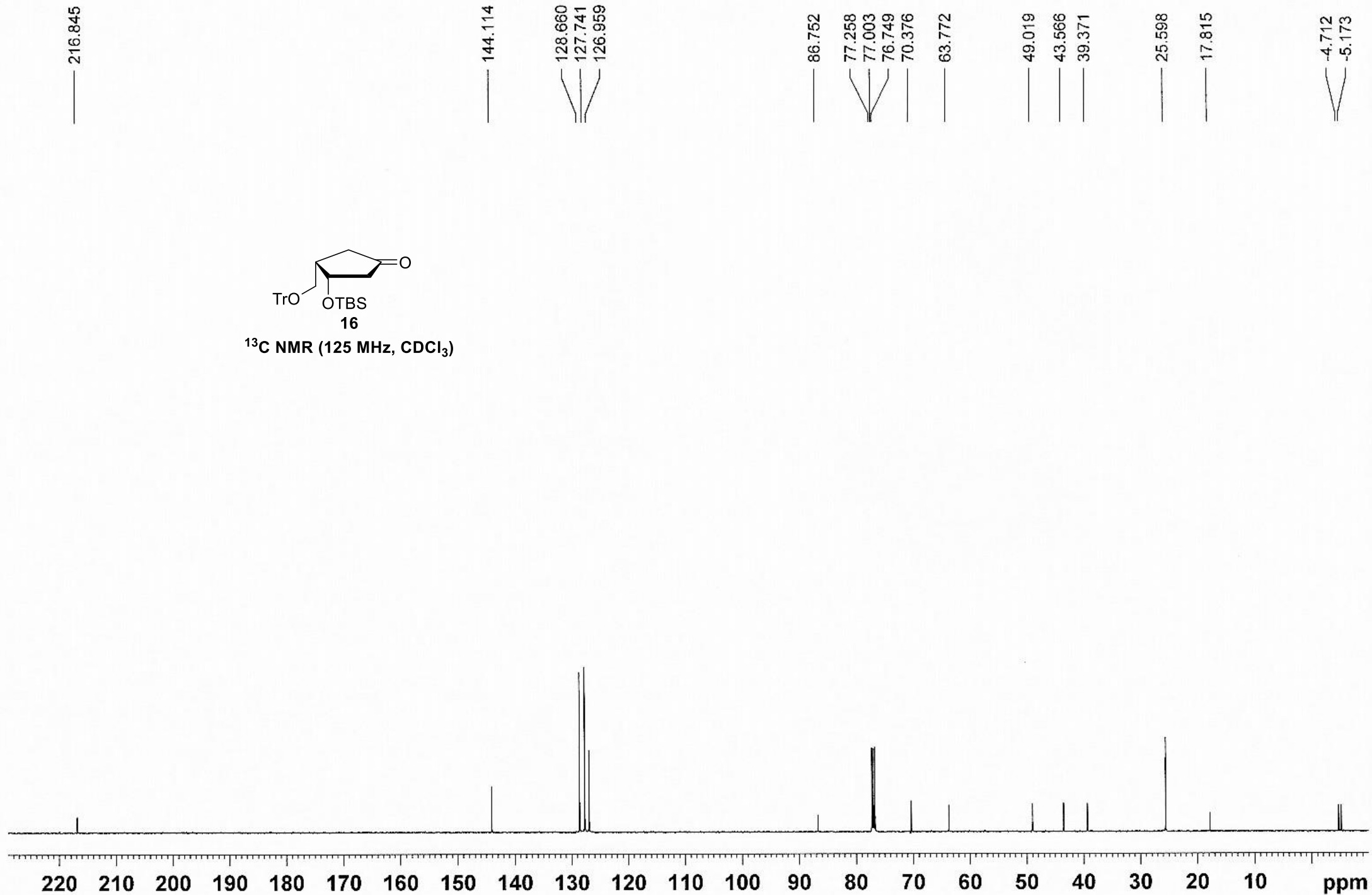
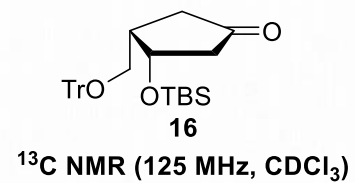
Elements : C 50/0, H 100/0, O 10/0

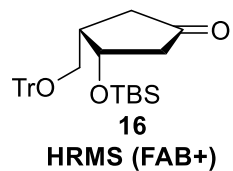
Mass Tolerance : 20ppm, 5mmu if m/z < 250, 10mmu if m/z > 500

Unsaturation (U.S.) : -0.5 - 20.0









Data : T0404005 Date : 04-Apr-2019 18:57

Instrument : MStation

Sample : B

Note : m-NBA

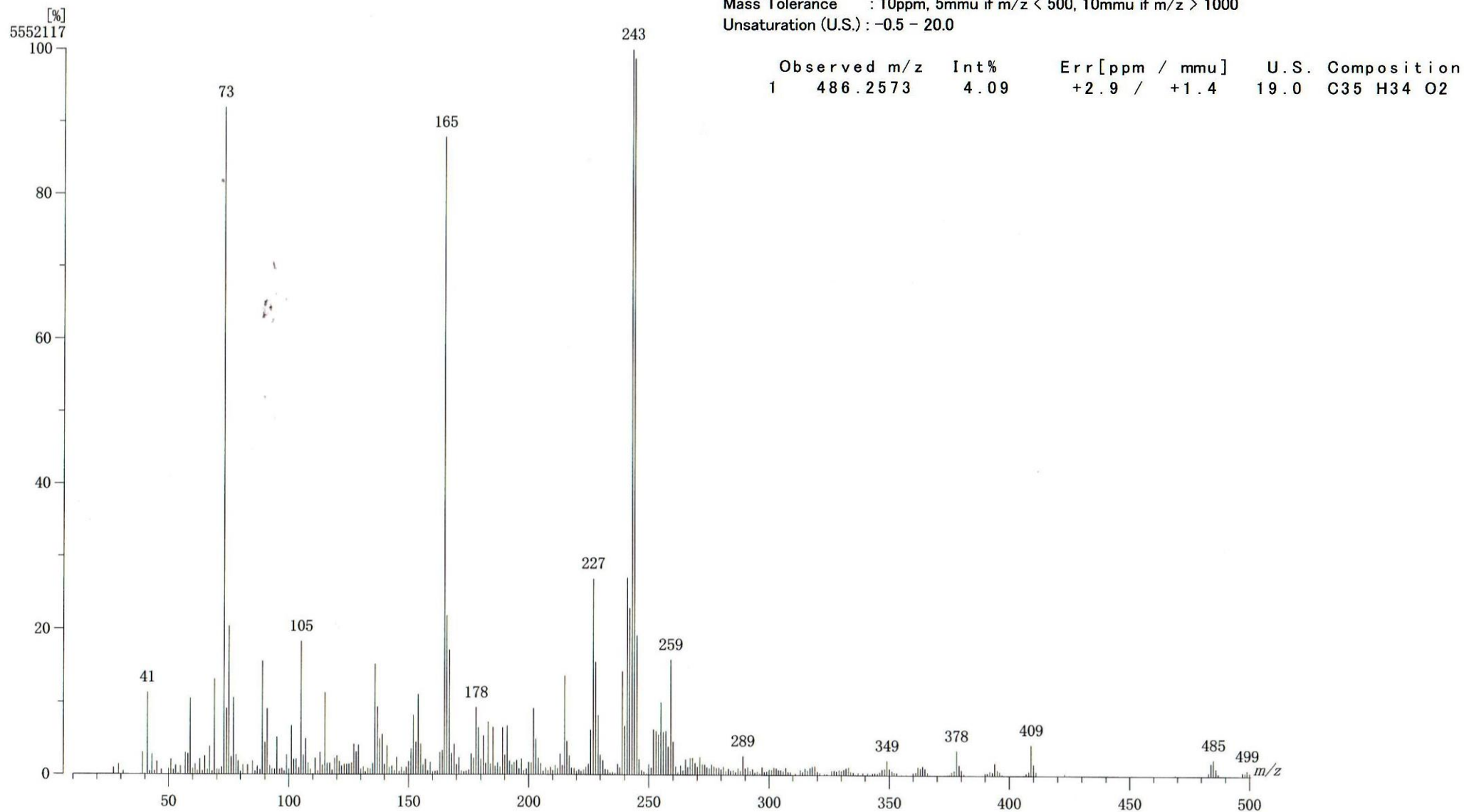
Inlet : Direct Ion Mode : FAB+

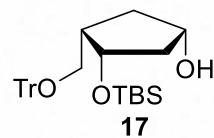
RT : 2.63 min Scan# : (46,52)

Elements : C 50/0, H 100/0, O 10/0, 28Si 1/0, 29Si 1/0, 30Si 1/0

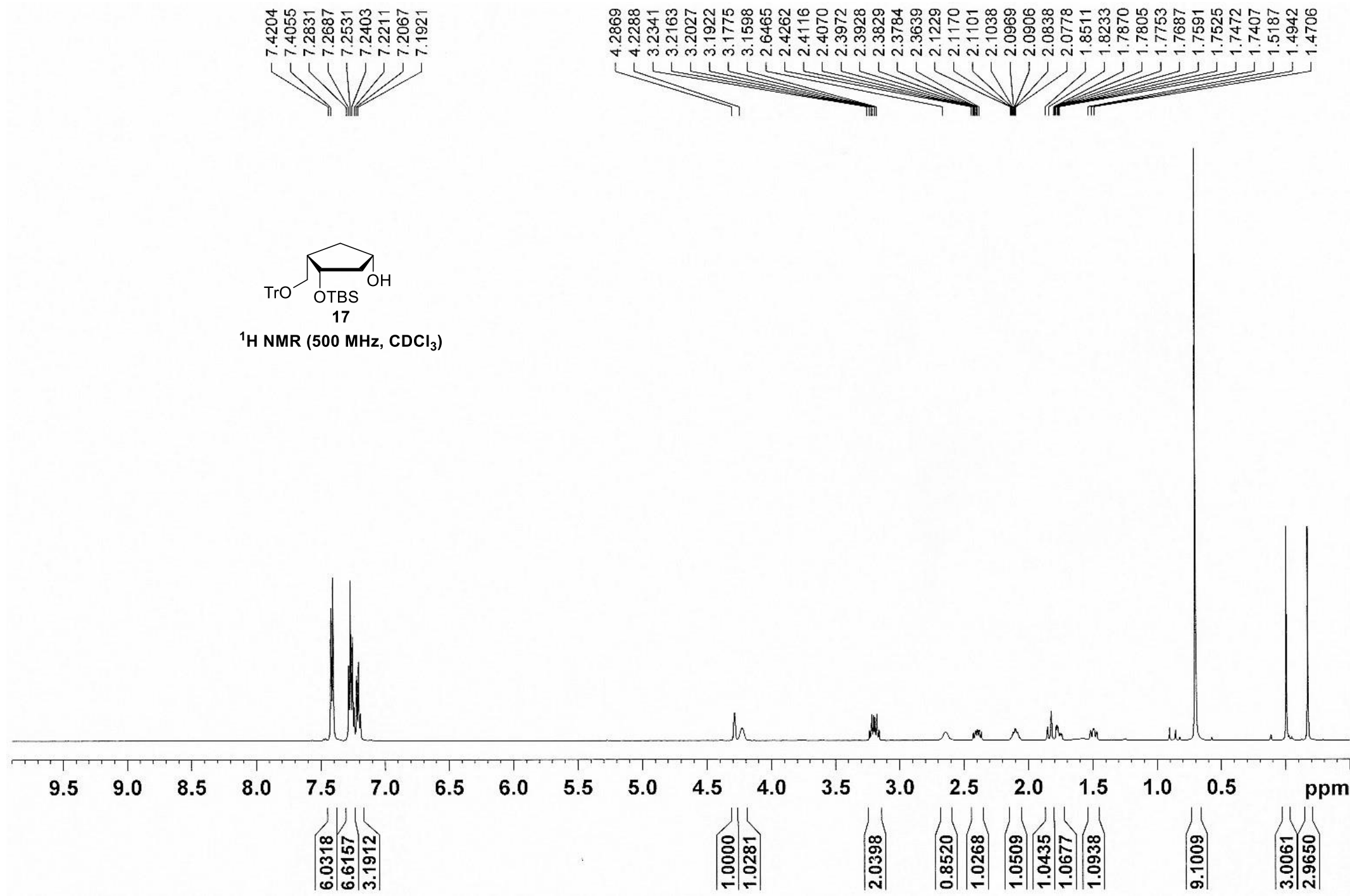
Mass Tolerance : 10ppm, 5mmu if m/z < 500, 10mmu if m/z > 1000

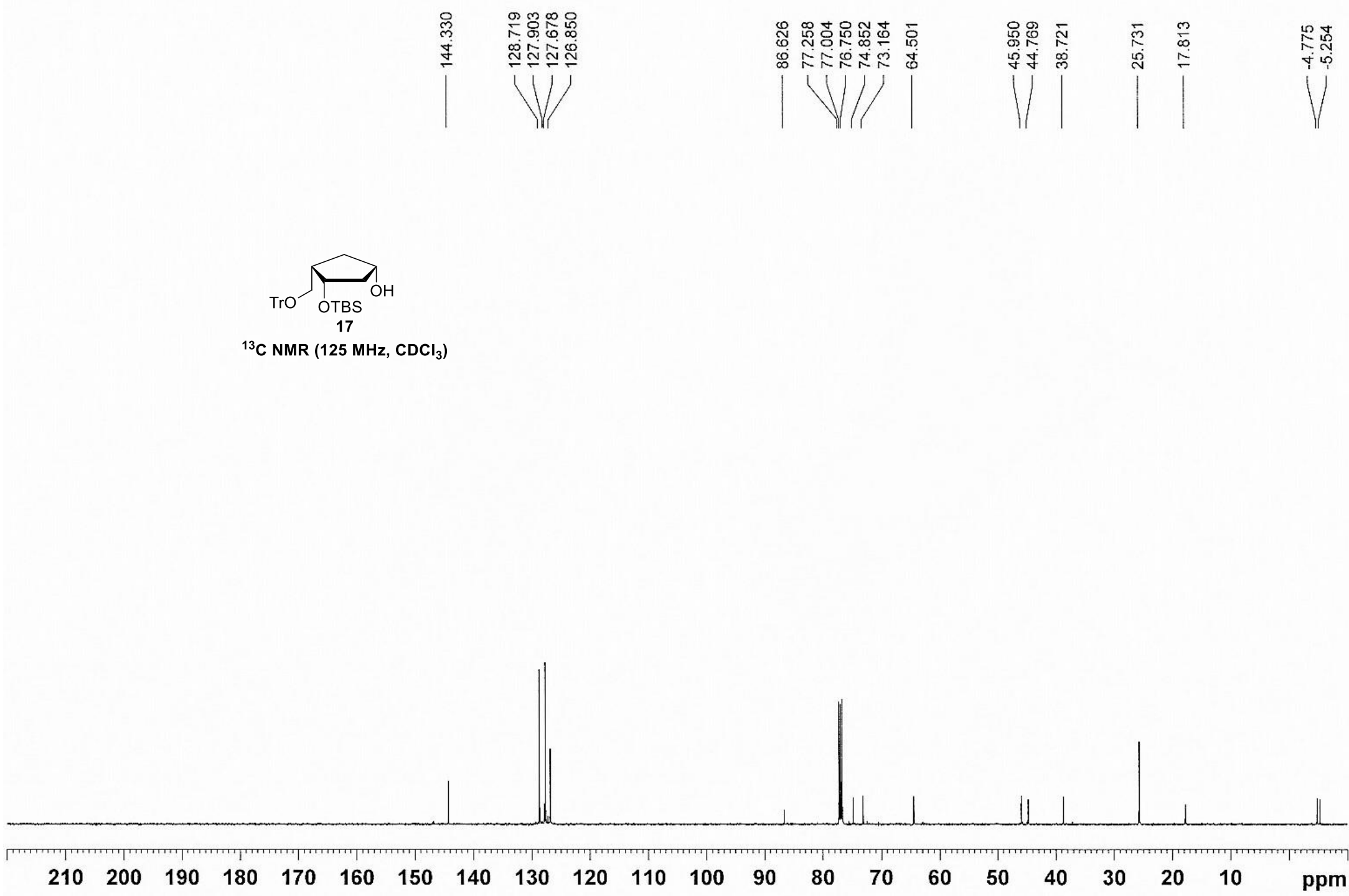
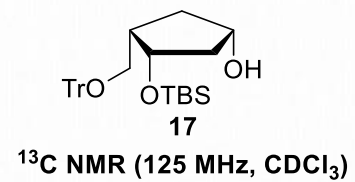
Unsaturation (U.S.) : -0.5 - 20.0





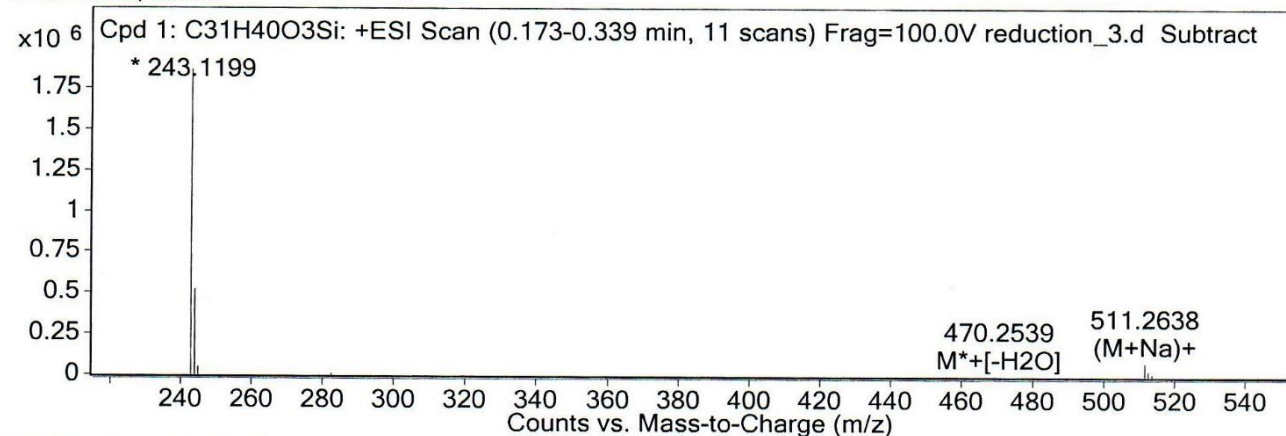
¹H NMR (500 MHz, CDCl₃)





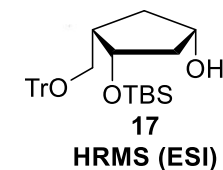
Qualitative Compound Report

MS Zoomed Spectrum

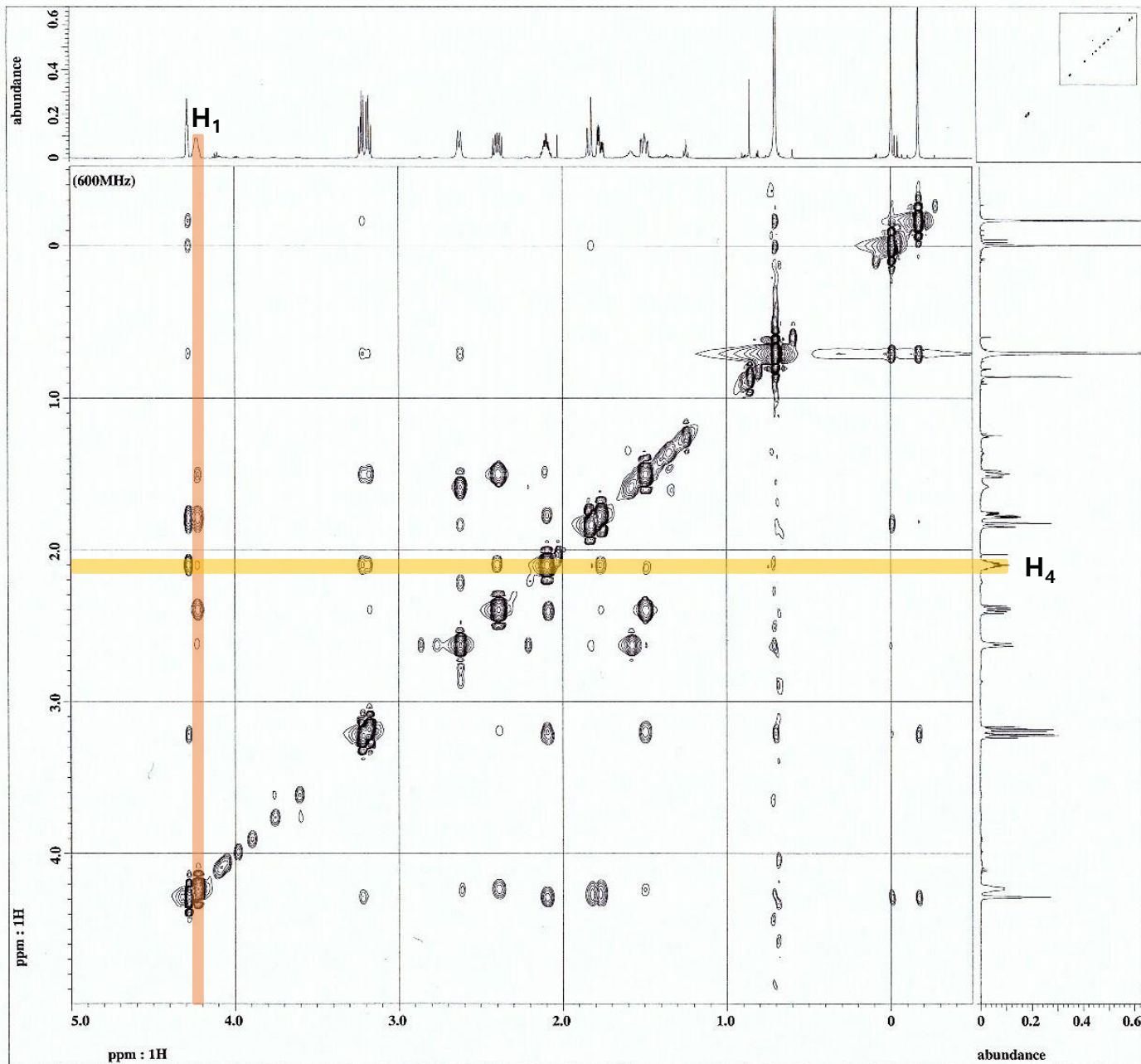
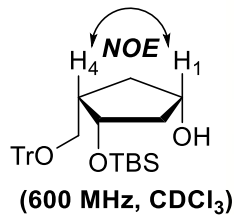


MS Spectrum Peak List

m/z	Calc m/z	Diff(ppm)	z	Abund	Formula	Ion
243.1199				1857543		
243.2565				116908		
243.2995				37826		
244.1195				526558		
244.1565				10165		
245.1241	245.1446	-83.86		45627	C ₃₁ H ₄₂ O ₃ Si	(M+2H)+2
470.2539	470.2636	-20.46	1	149	C ₃₁ H ₃₈ O ₂ Si	M*+[-H ₂ O]
511.2638	511.2639	-0.16	1	85309	C ₃₁ H ₄₀ Na O ₃ Si	(M+Na)+
512.2669	512.2668	0.12	1	32683	C ₃₁ H ₄₀ Na O ₃ Si	(M+Na)+
527.2372	527.2378	-1.28	1	1390	C ₃₁ H ₄₀ K O ₃ Si	(M+K)+



--- End Of Report ---



```

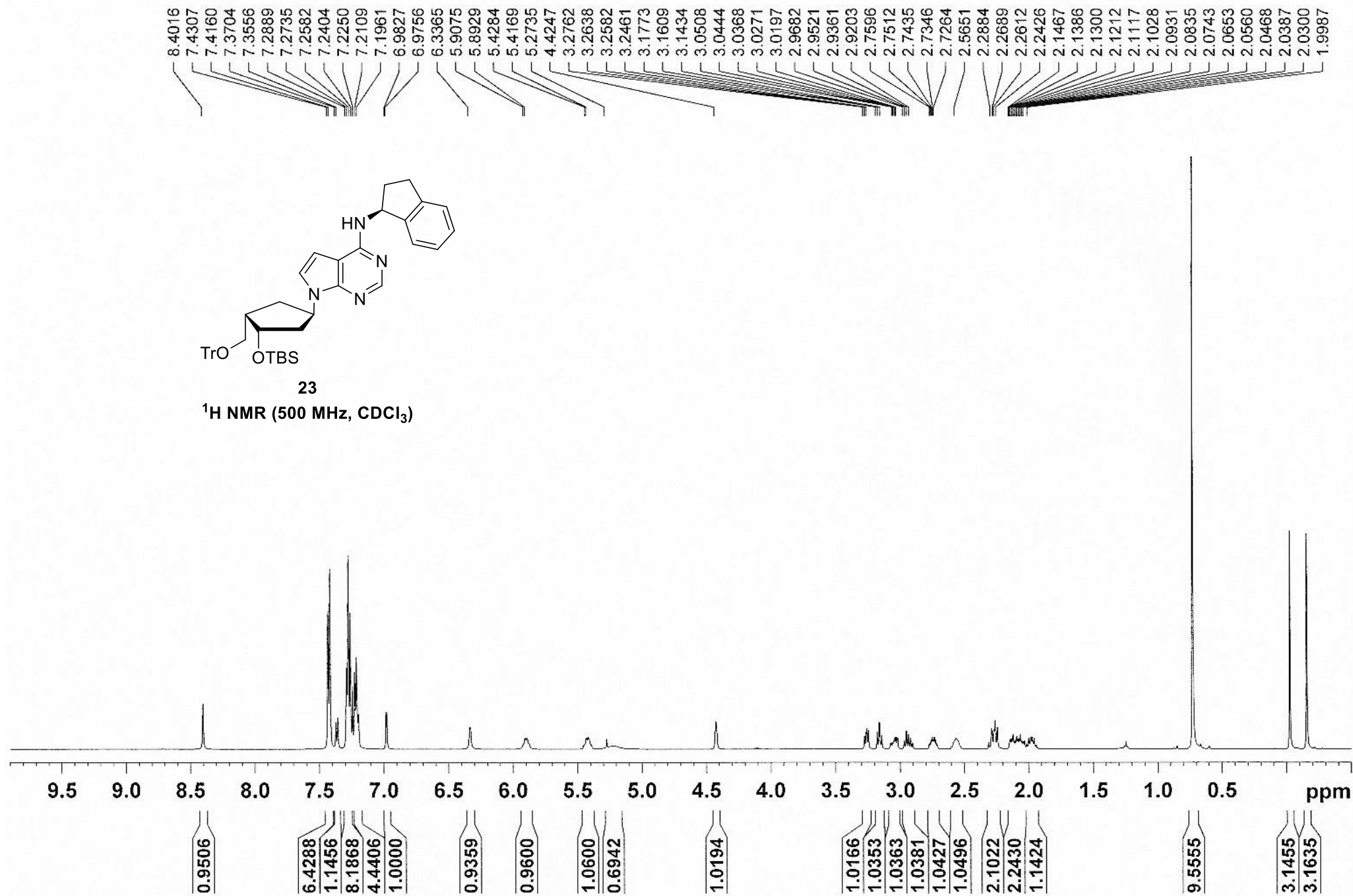
---- PROCESSING PARAMETERS ----
sexp : 5.0[Hz] : 0.0[s]
trapezoid3 : 0[%] : 80[%] : 100[%]
zerofill : 1
fft : 1 : TRUE : TRUE
ppm
machinephase
[transpose]
sexp : 5.0[Hz] : 0.0[s]
trapezoid3 : 0[%] : 80[%] : 100[%]
zerofill : 2
fft : 1 : TRUE : TRUE
ppm
machinephase

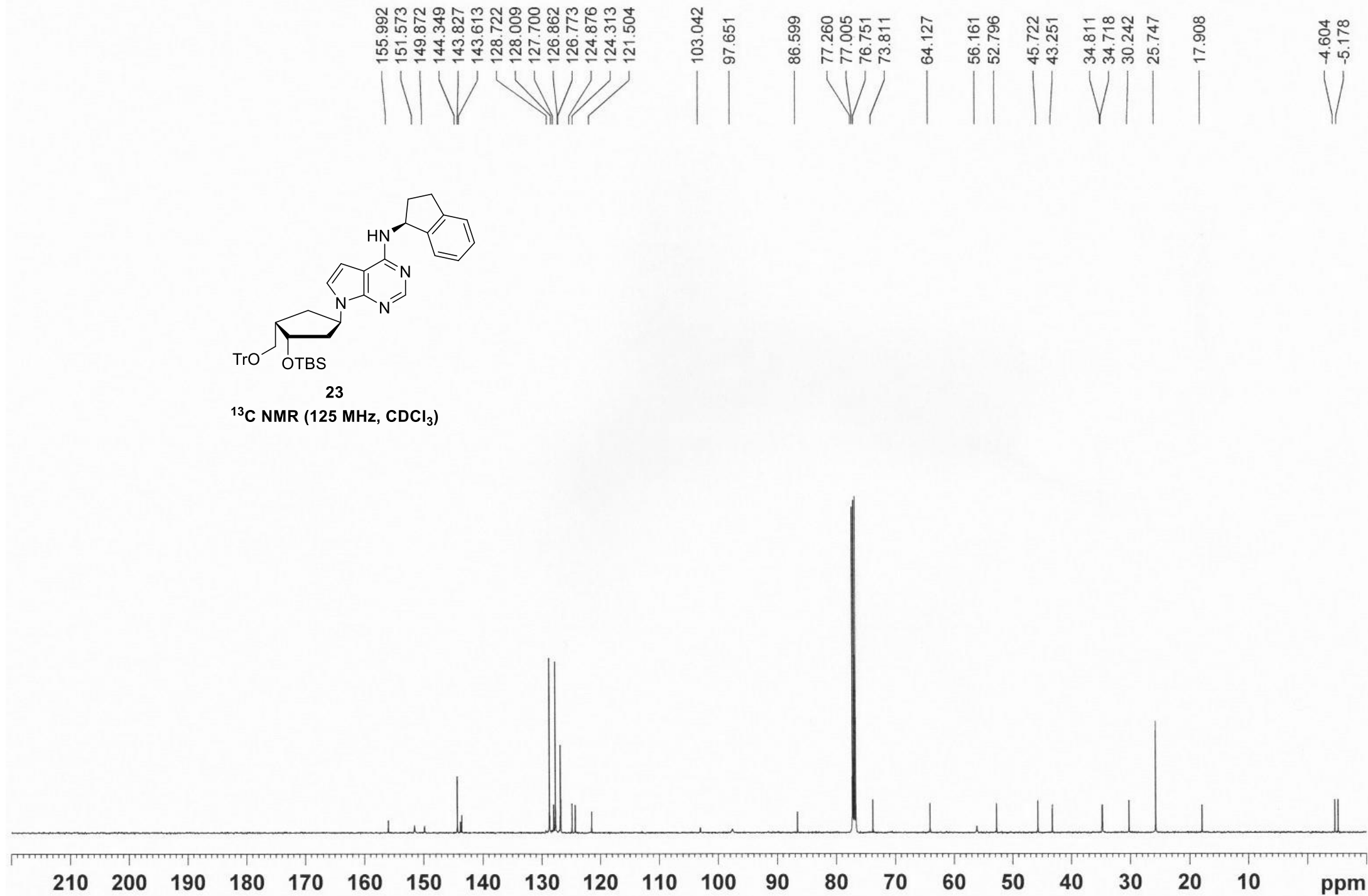
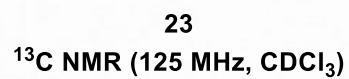
Filename      = NOESY-3.jdf
Author        = delta
Experiment    = noesy_phase_pfgz
Sample_id     = ft
Solvent       = CHLOROFORM-D
Creation_time  = 11-APR-2019 11:28:13
Revision_time = 11-APR-2019 16:41:42
Current_time  = 11-APR-2019 16:43:49

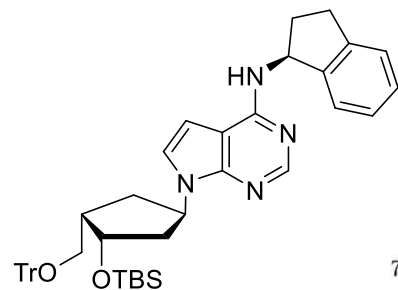
Comment       = (600MHz)
Data_format   = 2D REAL REAL
Dim_size      = 1638, 504
Dim_title     = 1M 1H
Dim_units     = [ppm] [ppm]
Dimensions    = X Y
Site          = ECA 600
Spectrometer  = JNM-ECA600

Field_strength = 14.09636928[T] (600[M
X_acq_duration = 0.22740992[s]
X_domain       = 1H
X_freq         = 600.1723046[MHz]
X_offset       = 5.00278[ppm]
X_points       = 2048
X_prescans     = 4
X_resolution   = 4.39734555[Hz]
X_sweep        = 9.00576369[kHz]
Y_domain       = 1H
Y_freq         = 600.1723046[MHz]
Y_offset       = 5.00278[ppm]
Y_points       = 252
Y_prescans     = 0
Y_resolution   = 28.58148926[Hz]
Y_sweep        = 7.20253529[kHz]
Irr_domain     = 1H
Irr_freq       = 600.1723046[MHz]
Irr_offset     = 5[ppm]
Tri_domain     = 1H
Tri_freq       = 600.1723046[MHz]
Tri_offset     = 5[ppm]
Clipped        = FALSE
Mod_return     = 1
Scans          = 12
Total_scans    = 3024

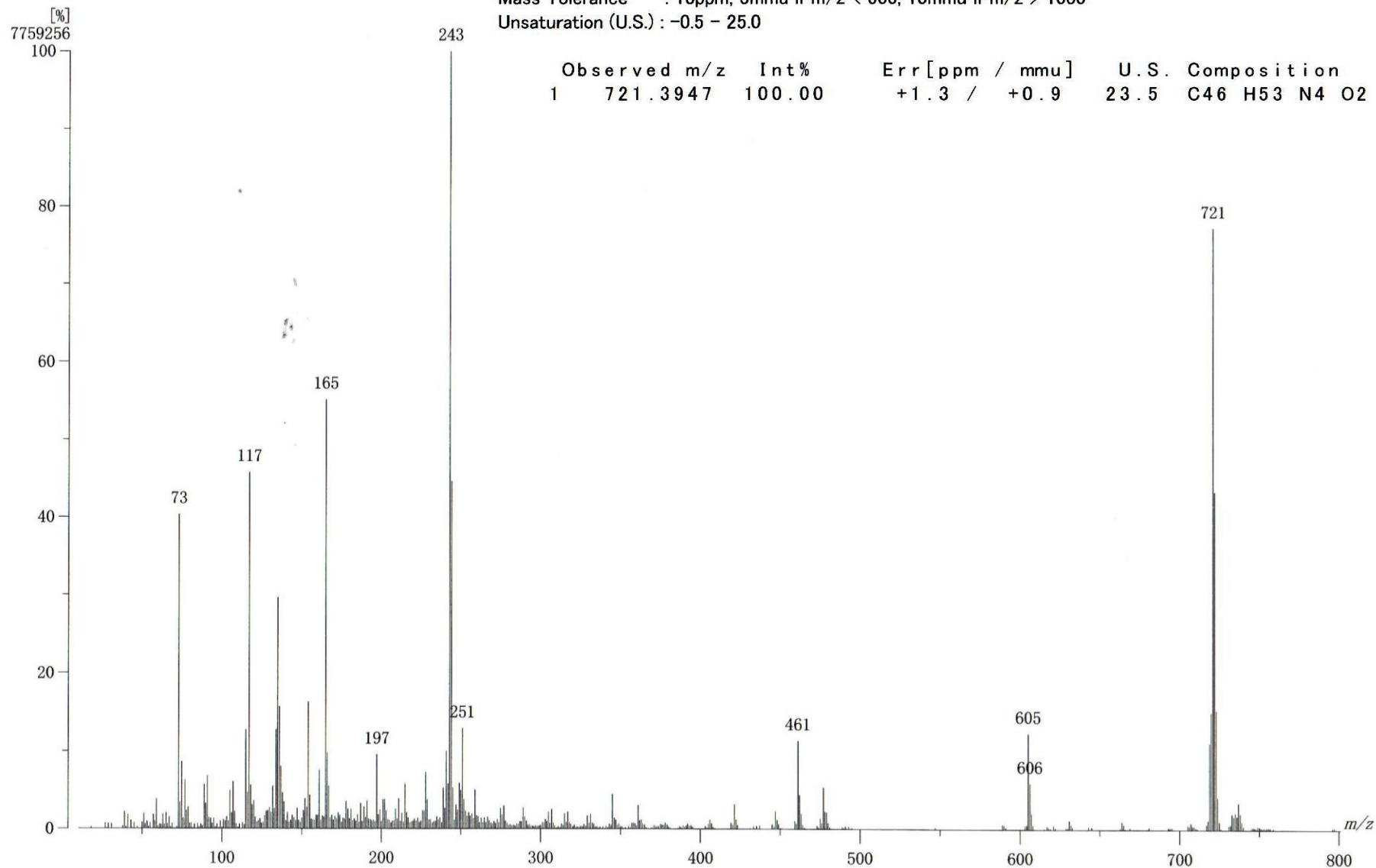
X_acq_time     = 0.22740992[s]
X_atn          = 4.4[dB]
X_pulse        = 6.36[us]
Y_acq_time     = 35.54304[ms]
Y_pl_correction = 180
Irr_mode       = Off
Tri_mode       = Off
Dante_presat   = FALSE
Grad_1         = 1[ms]
Grad_1_amp     = 45[mT/m]
Initial_wait   = 1[s]
Mix_time       = 0.5[s]
Recvr_gain     = 34
Relaxation_delay = 1.5[s]
Repetition_time = 1.72740992[s]
  
```







23
HRMS (FAB+)



Data : T0404008 Date : 05-Apr-2019 09:42

Instrument : MStation

Sample : D

Note : m-NBA

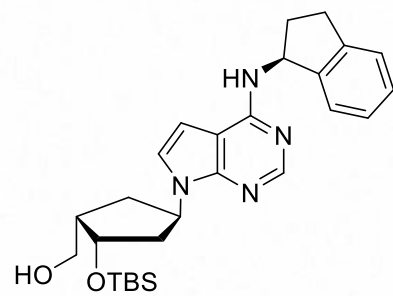
Inlet : Direct Ion Mode : FAB+

RT : 0.00 min Scan# : (1,59)

Elements : C 50/0, H 100/0, N 4/0, O 3/0, 28Si 1/0, 29Si 1/0, 30Si 1/0

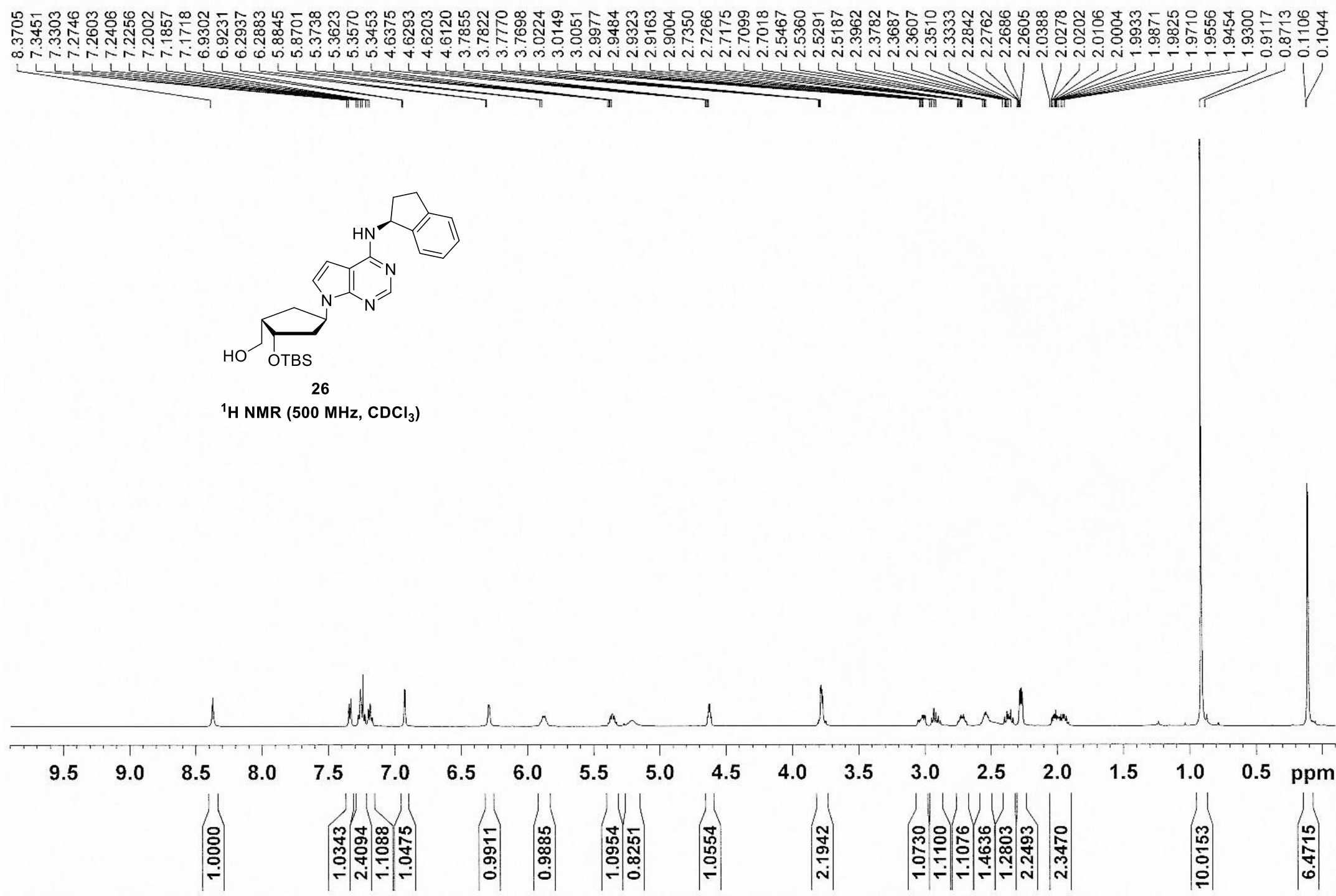
Mass Tolerance : 10ppm, 5mmu if m/z < 500, 10mmu if m/z > 1000

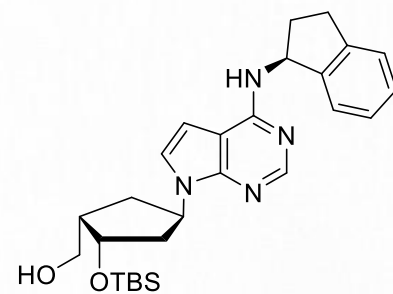
Unsaturation (U.S.) : -0.5 - 25.0



26

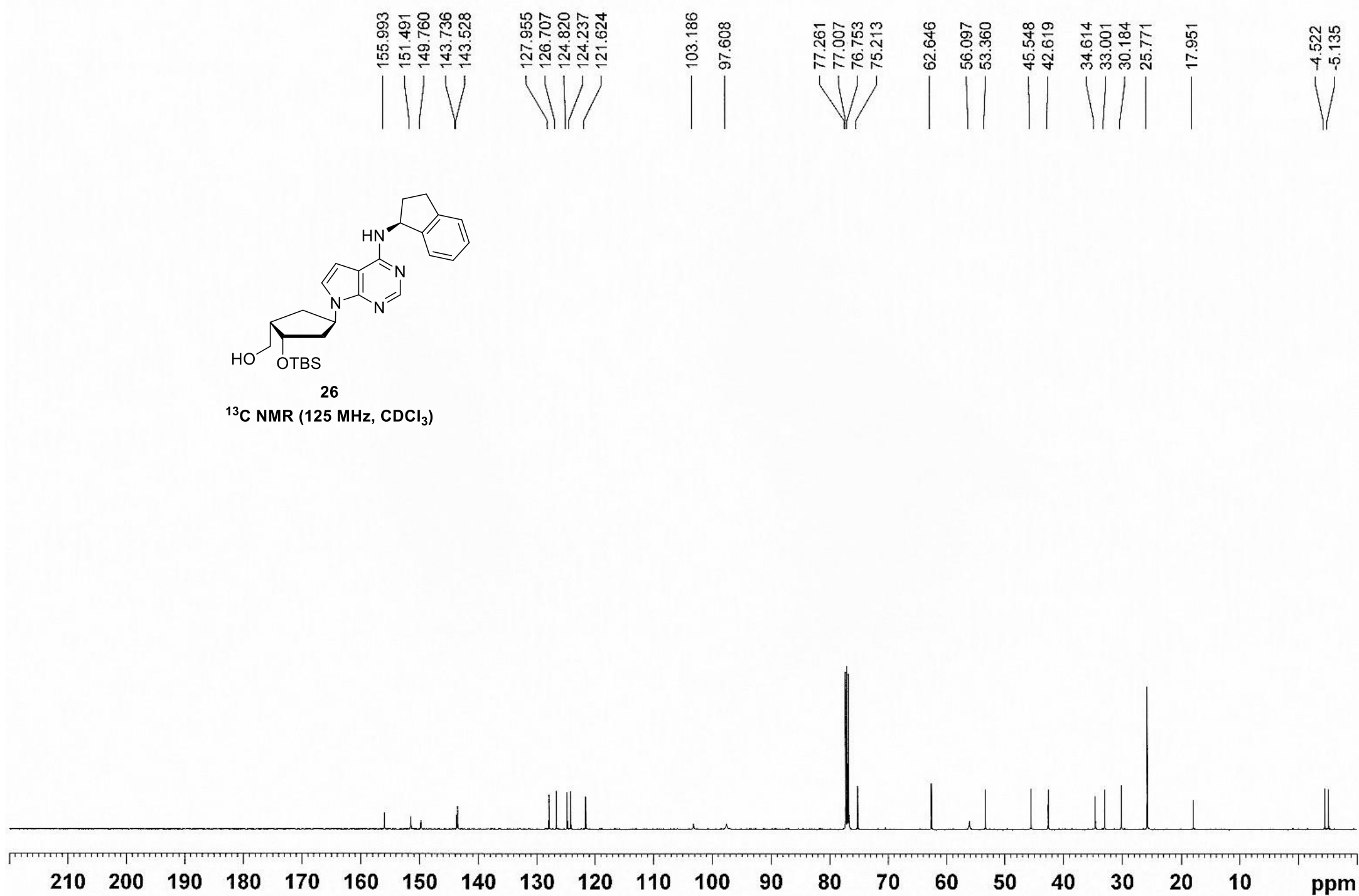
^1H NMR (500 MHz, CDCl_3)

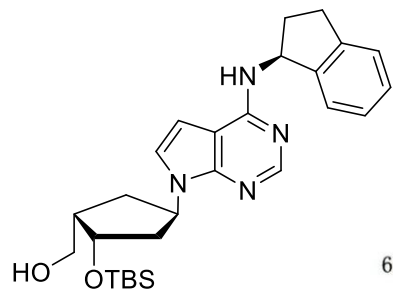




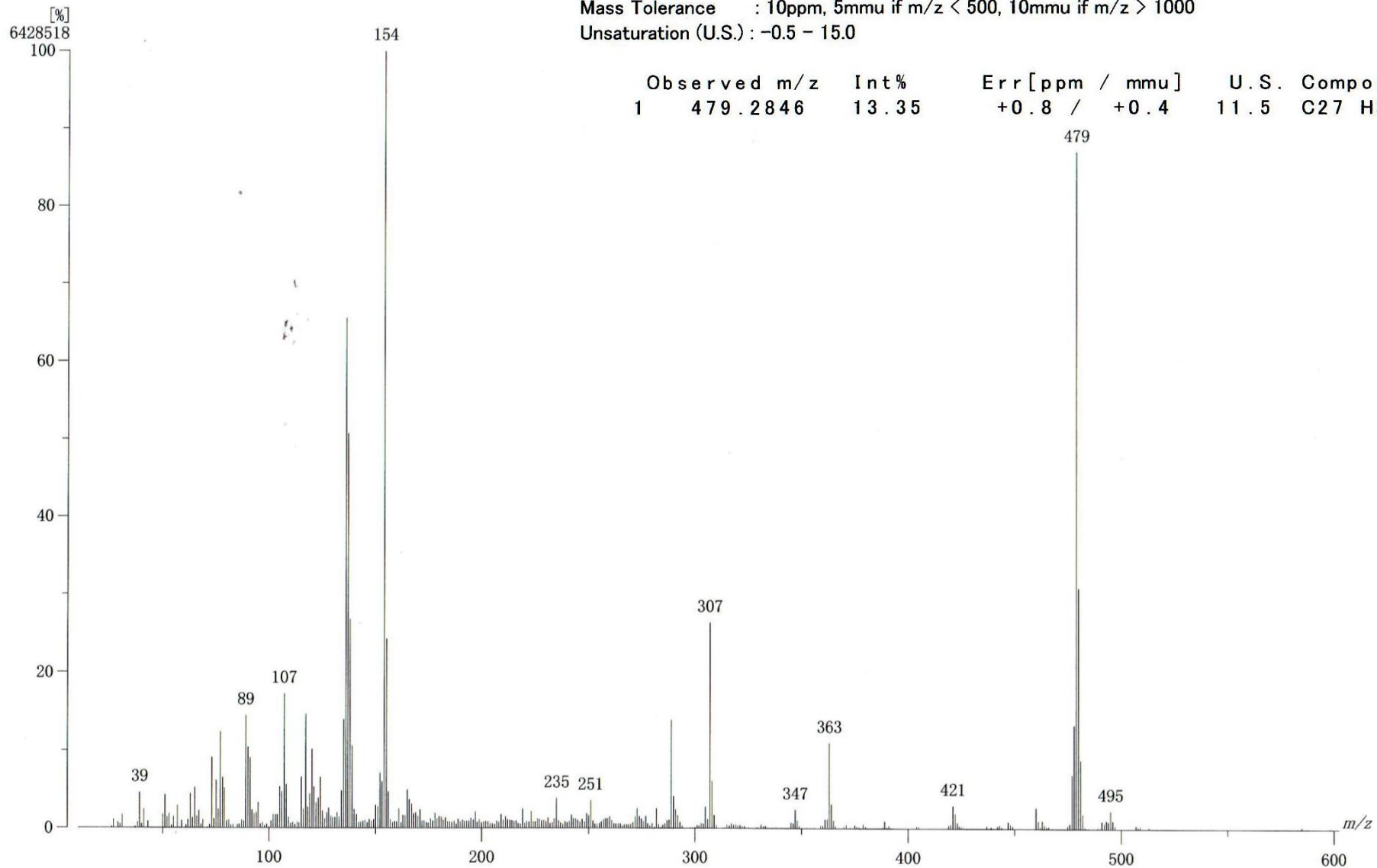
26

^{13}C NMR (125 MHz, CDCl_3)





26
HRMS (FAB+)



Data : T0404010 Date : 05-Apr-2019 11:33

Instrument : MStation

Sample : E

Note : m-NBA

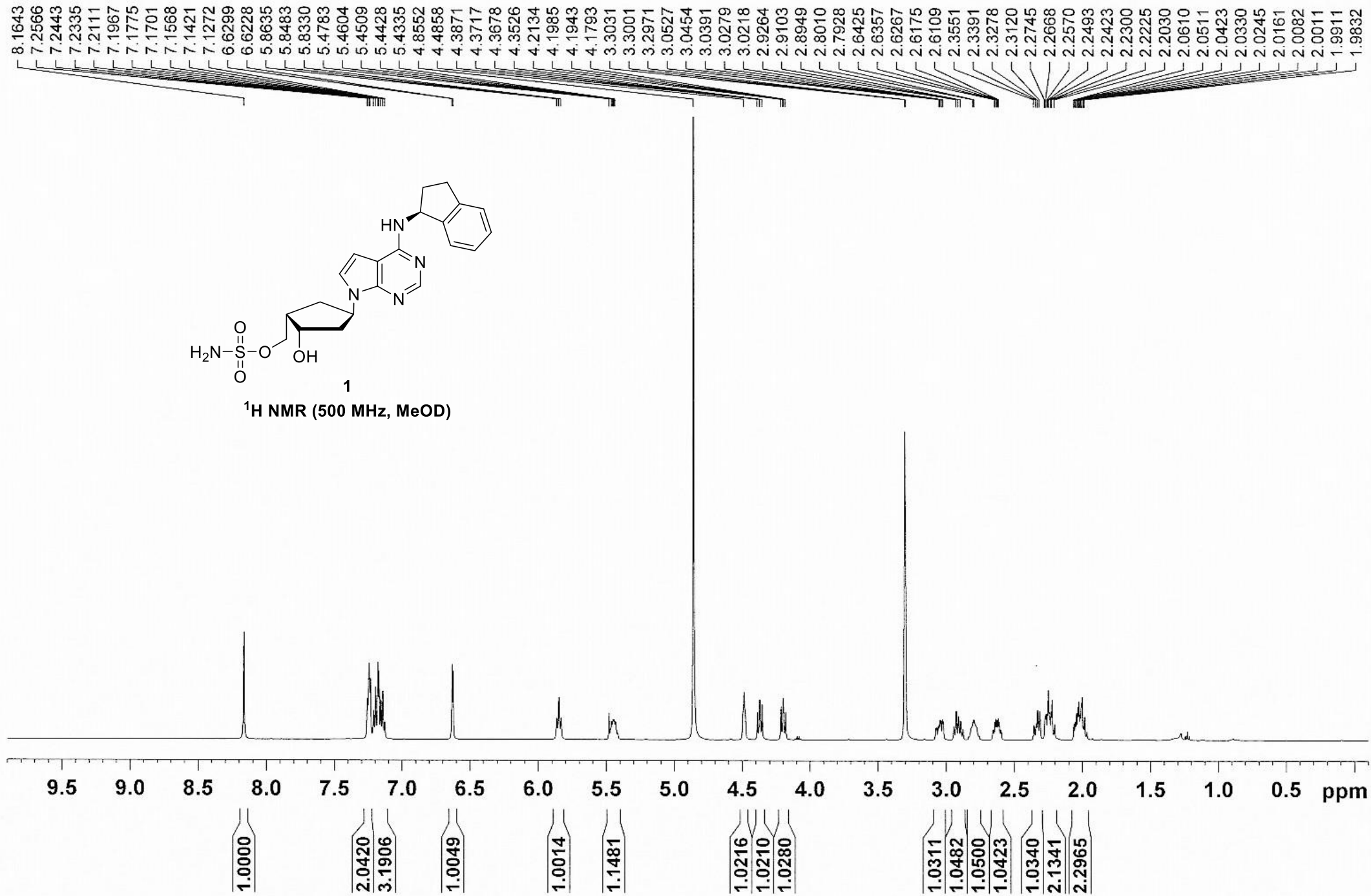
Inlet : Direct Ion Mode : FAB+

RT : 1.87 min Scan# : (33,57)

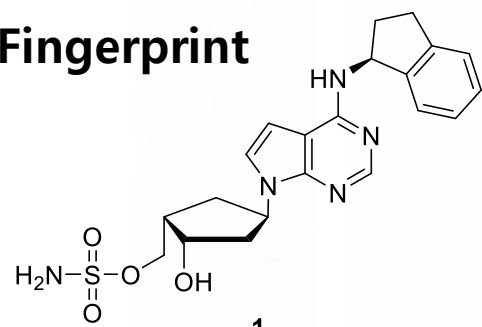
Elements : C 50/0, H 100/0, N 4/0, O 3/0, 28Si 1/0, 29Si 1/0, 30Si 1/0

Mass Tolerance : 10ppm, 5mmu if m/z < 500, 10mmu if m/z > 1000

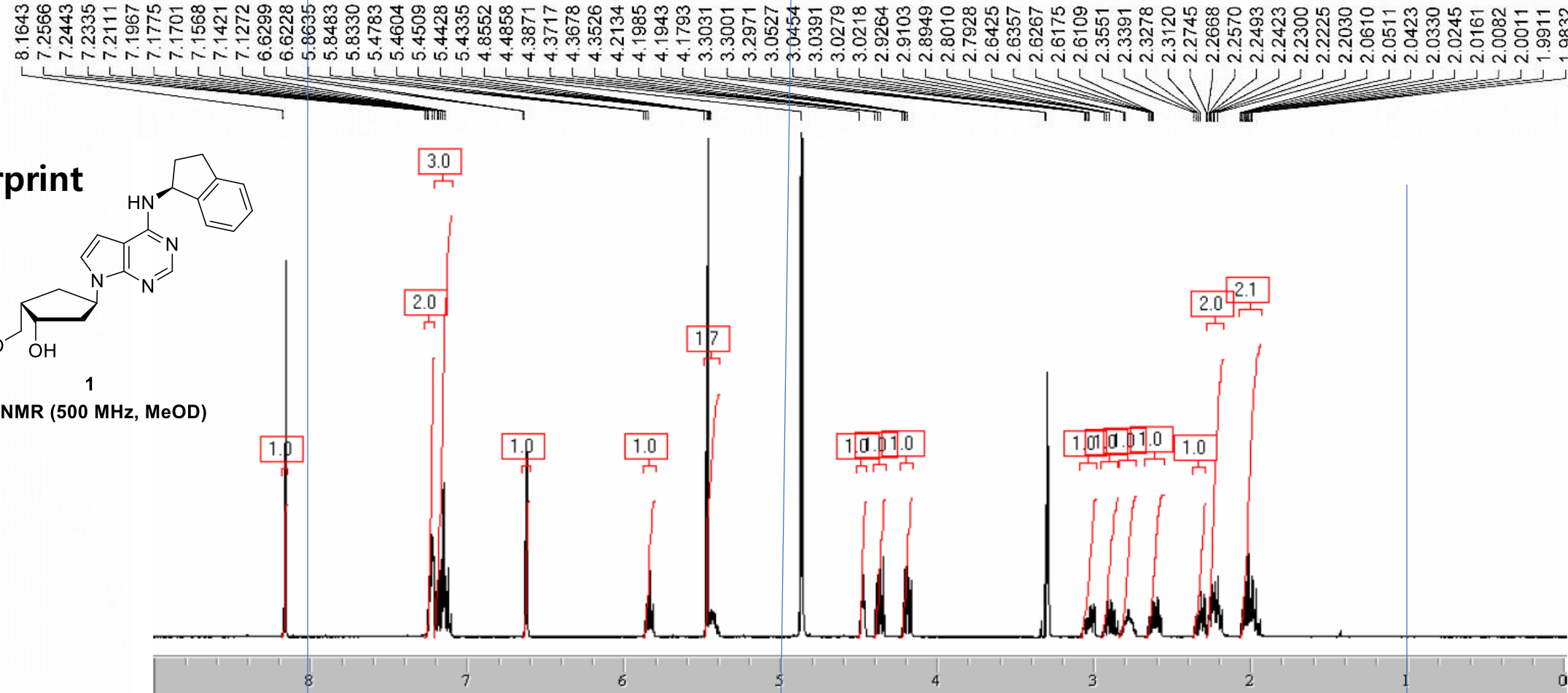
Unsaturation (U.S.) : -0.5 - 15.0



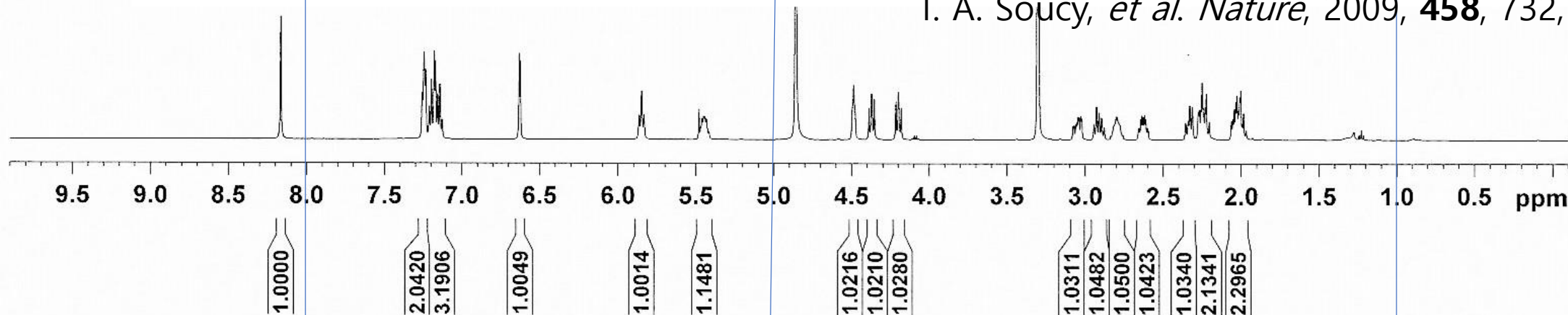
Fingerprint

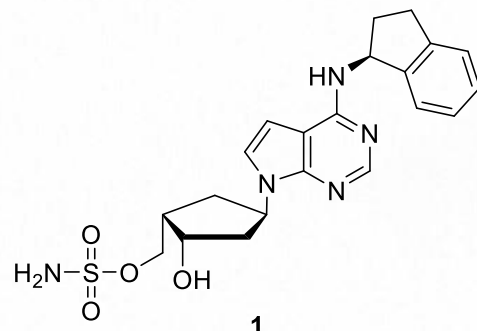


¹H NMR (500 MHz, MeOD)

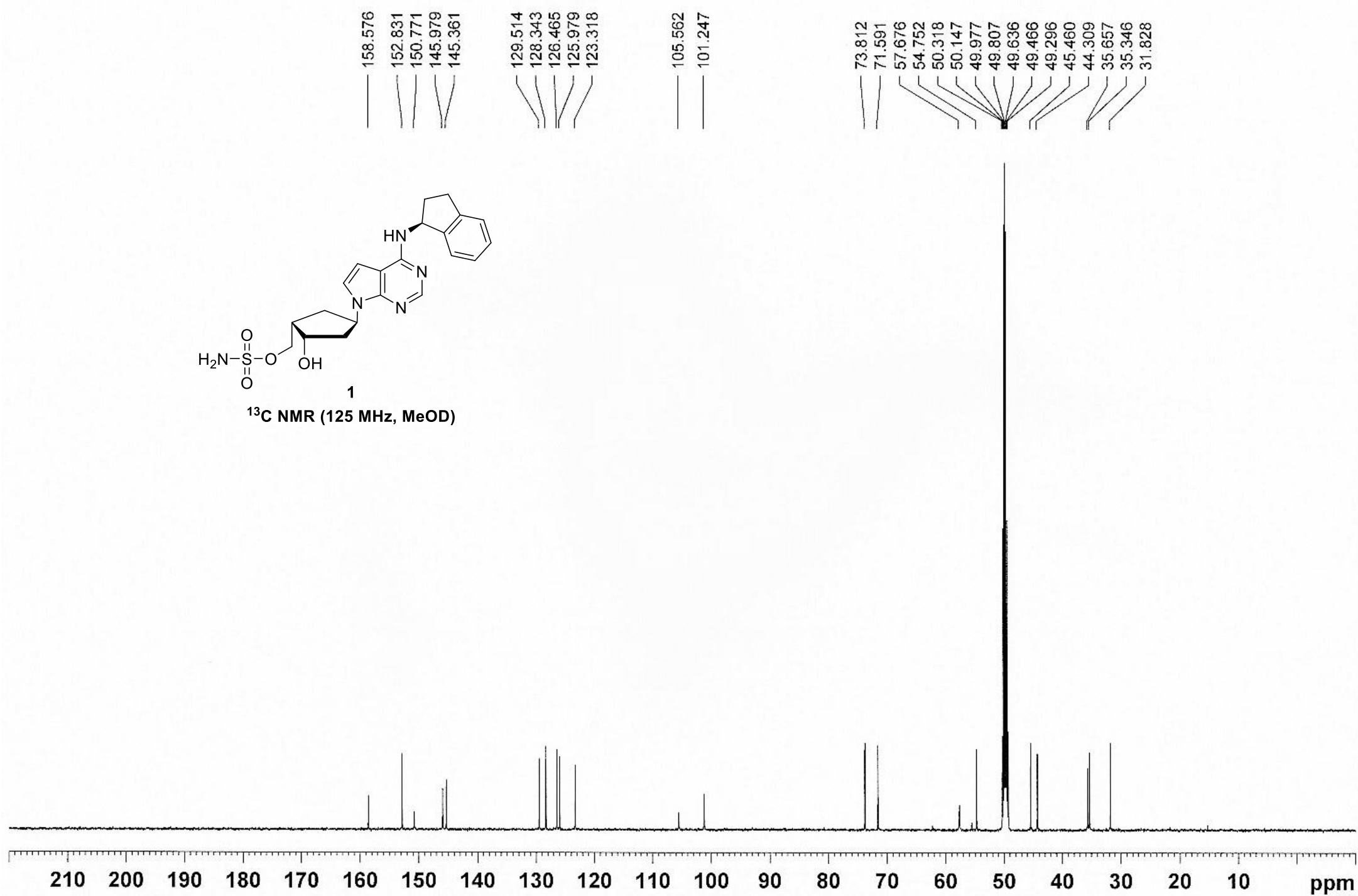


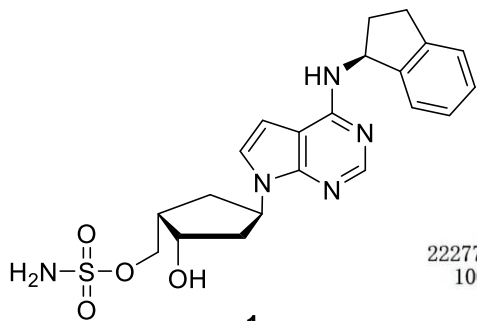
T. A. Soucy, *et al.* *Nature*, 2009, **458**, 732,



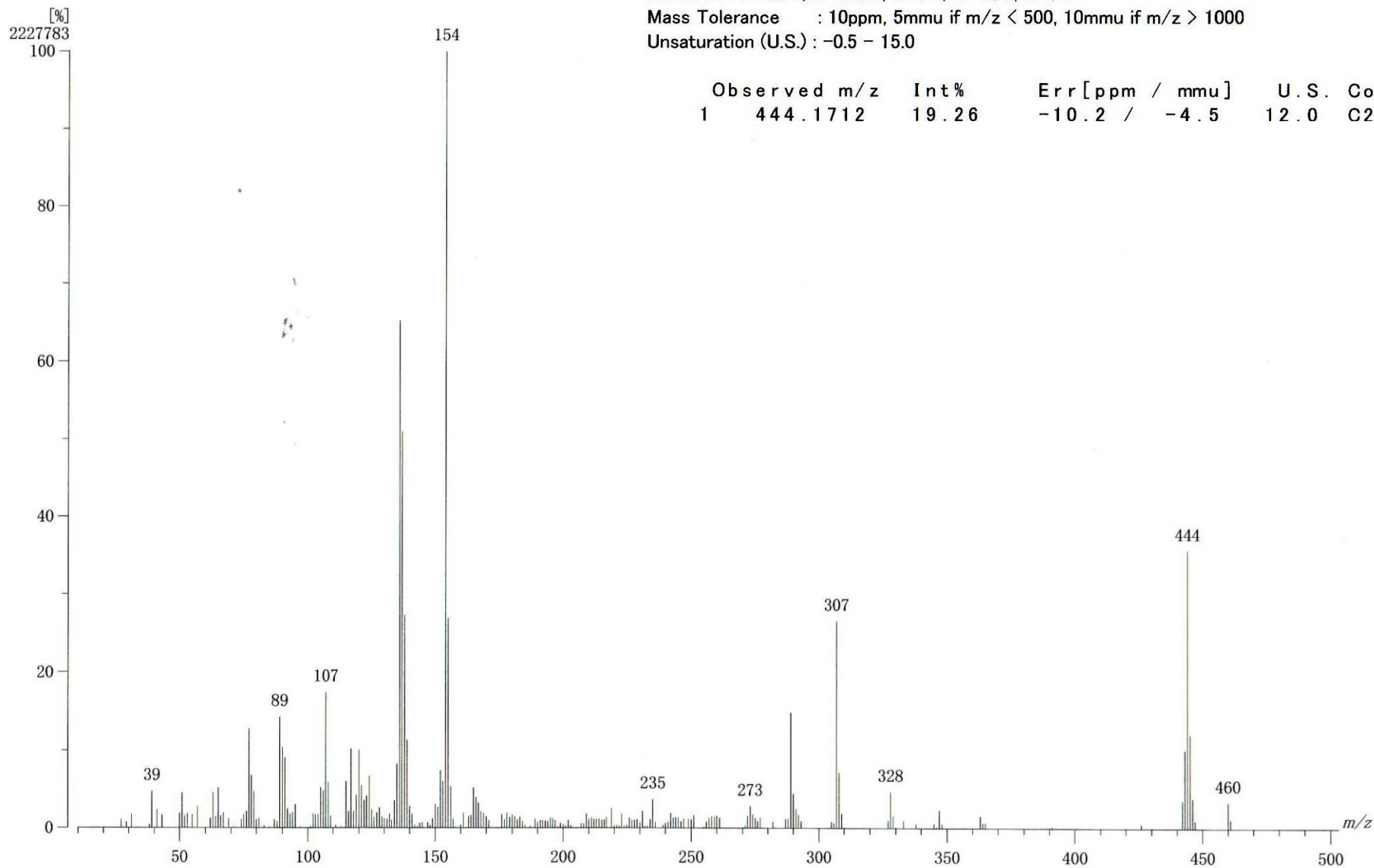


¹³C NMR (125 MHz, MeOD)





1
HRMS (FAB+)



Data : T0404012 Date : 05-Apr-2019 14:13

Instrument : MStation

Sample : F

Note : m-NBA

Inlet : Direct Ion Mode : FAB+

RT : 0.00 min Scan# : (1,42)

Elements : C 50/0, H 100/0, N 6/0, O 10/0, S 1/0

Mass Tolerance : 10ppm, 5mmu if m/z < 500, 10mmu if m/z > 1000

Unsaturation (U.S.) : -0.5 - 15.0

	Observed m/z	Int%	Err[ppm / mmu]	U.S. Composition
1	444.1712	19.26	-10.2 / -4.5	12.0 C20 H24 N6 O6

