

Supporting Information

Synthesis and Evaluation of Eight- and Four-membered Iminosugar Analogues as Inhibitors of Testicular Ceramide-specific Glucosyltransferase, Testicular β -Glucosidase 2, and other Glycosidases

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REFERENCE NUMBER: 08001

CRYSTAL STRUCTURE REPORT

$2(\text{C}_{16}\text{H}_{33}\text{N}\text{O}_5); \text{H}_2\text{O}$

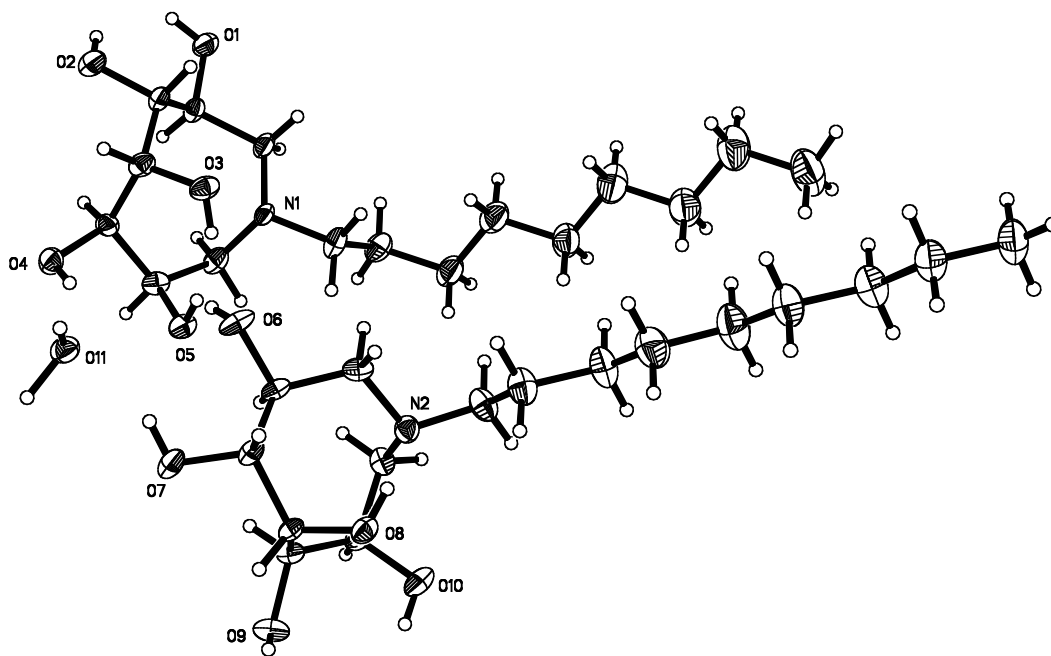
$\text{C}_{32}\text{H}_{68}\text{N}_2\text{O}_{11}$

Report prepared for:

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Data collection

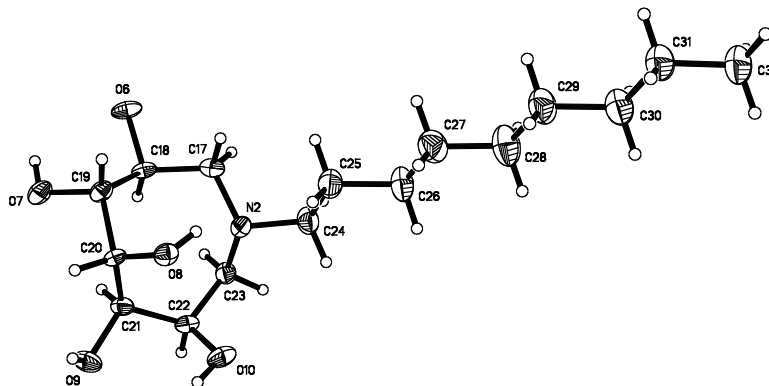
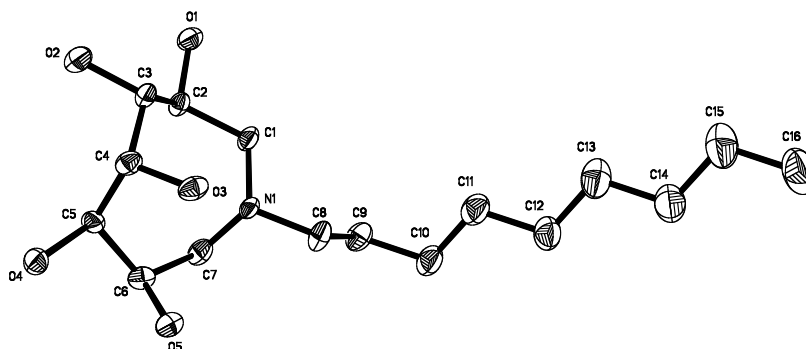
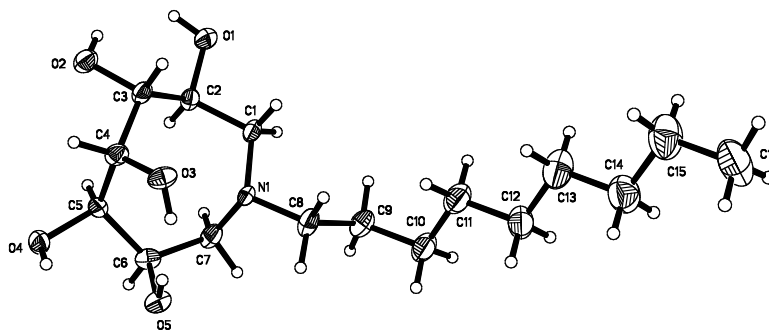
A crystal (approximate dimensions $0.35 \times 0.30 \times 0.02 \text{ mm}^3$) was placed onto the tip of a 0.1 mm diameter glass capillary and mounted on a Bruker SMART Platform CCD diffractometer for a data collection at 173(2) K.¹ A preliminary set of cell constants was calculated from reflections harvested from three sets of 20 frames. These initial sets of frames were oriented such that orthogonal wedges of reciprocal space were surveyed. This produced initial orientation matrices determined from 93 reflections. The data collection was carried out using MoK α radiation (graphite monochromator) with a frame time of 120 seconds and a detector distance of 4.86 cm. A randomly oriented region of reciprocal space was surveyed to the extent of one sphere and to a resolution of 0.84 Å. Four major sections of frames were collected with 0.30° steps in ω at four different ϕ settings and a detector position of -28° in 2θ . The intensity data were corrected for absorption and decay (SADABS).² Final cell constants were calculated from the xyz centroids of 2524 strong reflections from the actual data collection after integration (SAINT).³ Please refer to Table 1 for additional crystal and refinement information.

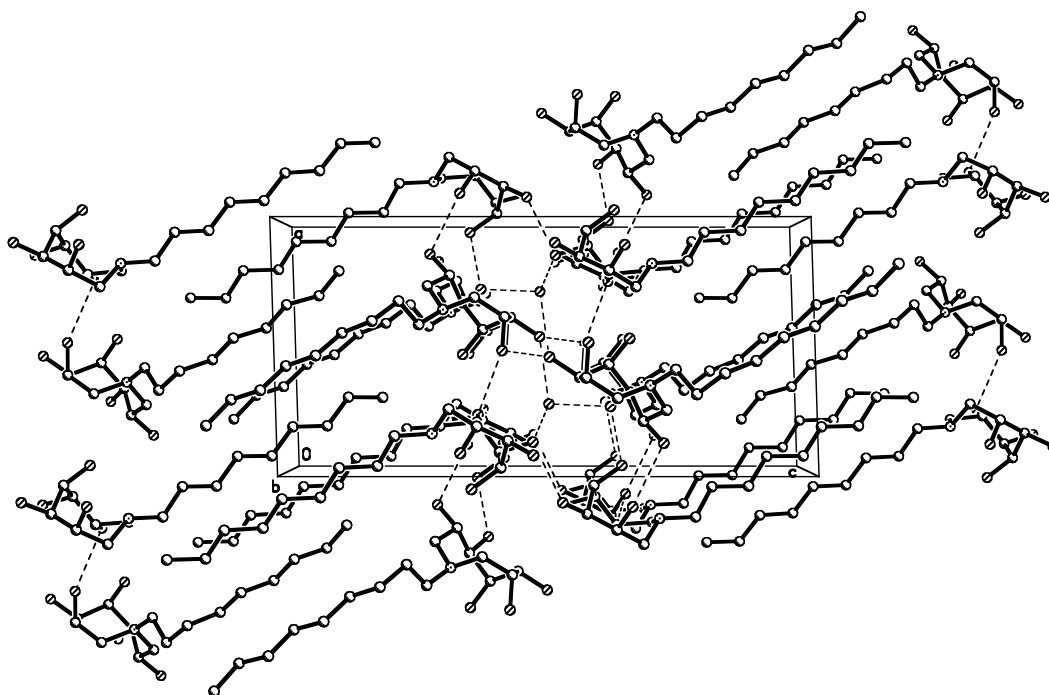
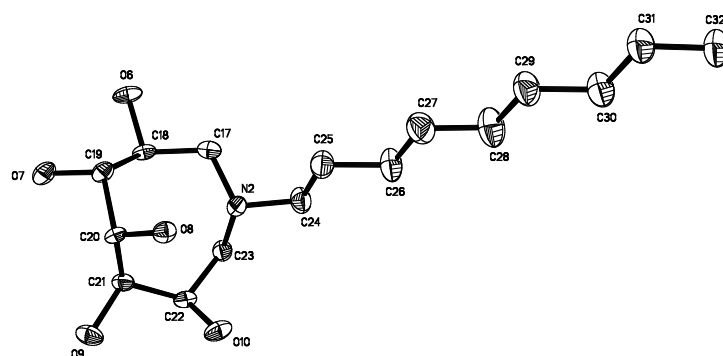
Structure solution and refinement

The structure was solved using SHELXS-97⁴ and refined using SHELXL-97.⁴ The space group $P2_1$ was determined based on systematic absences and intensity statistics. A direct-methods solution was calculated which provided most non-hydrogen atoms from the E-map. Full-matrix least squares / difference Fourier cycles were performed which located the remaining non-hydrogen atoms. All non-hydrogen atoms were refined with anisotropic displacement parameters. All hydrogen atoms were placed in ideal positions and refined as riding atoms with relative isotropic displacement parameters. The final full matrix least squares refinement converged to $R1 = 0.0527$ and $wR2 = 0.1700$ (F^2 , all data).

Structure description

The structure is the one suggested. There are two identical molecules in the asymmetric unit, as well as one water molecule. The absolute stereochemistry cannot be assigned from the diffraction data due to the lack of a heavy atom ($> \text{Si}$); however, the chemist has assigned the stereochemistry based on a known chiral center. The stereochemistry of C2, C3, C5 and C6 are S, R, R, and R respectively; and the stereochemistry of C18, C19, C21 and C22 are S, R, R, and R respectively. The two molecules have several hydrogen bond interactions to each other, as well as to the water molecule.





Matrix Packing Plot (looking down the b-axis, hydrogen atoms omitted for clarity)

Data collection and structure solution were conducted at the X-Ray Crystallographic Laboratory, S146 Kolthoff Hall, Department of Chemistry, University of Minnesota. All calculations were performed using Pentium computers using the current SHELXTL suite of programs. All publications arising from this report MUST either 1)include Benjamin E. Kucera as a coauthor or 2)acknowledge Benjamin E. Kucera, Victor G. Young, Jr., and the X-Ray Crystallographic Laboratory.

References:

1. SMART V5.054, Bruker Analytical X-ray Systems, Madison, WI (2001).
2. An empirical correction for absorption anisotropy, R. Blessing, *Acta Cryst.* **A51**, 33-38 (1995).
3. SAINT+ V6.45, Bruker Analytical X-Ray Systems, Madison, WI (2003).
4. SHELXTL V6.14, Bruker Analytical X-Ray Systems, Madison, WI (2000).

Some equations of interest:

$$R_{\text{int}} = \Sigma |F_o^2 - \langle F_o^2 \rangle| / \Sigma |F_o^2|$$

$$R_1 = \Sigma ||F_o| - |F_c|| / \Sigma |F_o|$$

$$wR2 = [\Sigma [w(F_o^2 - F_c^2)^2] / \Sigma [w(F_o^2)^2]]^{1/2}$$

$$\text{where } w = q / [\sigma^2 (F_o^2) + (a*P)^2 + b*P + d + e*\sin(\theta)]$$

$$\text{Goof} = S = [\Sigma [w(F_o^2 - F_c^2)^2] / (n-p)]^{1/2}$$

Table 1. Crystal data and structure refinement for 08001.

Identification code	08001	
Empirical formula	$C_{32} H_{68} N_2 O_{11}$	
Formula weight	656.88	
Temperature	173(2) K	
Wavelength	0.71073 Å	
Crystal system	Monoclinic	
Space group	$P2_1$	
Unit cell dimensions	$a = 10.3632(13)$ Å	$\alpha = 90^\circ$
	$b = 8.1773(10)$ Å	$\beta = 91.751(2)^\circ$
	$c = 21.547(3)$ Å	$\gamma = 90^\circ$
Volume	$1825.1(4)$ Å ³	
Z	2	
Density (calculated)	1.195 Mg/m ³	
Absorption coefficient	0.088 mm ⁻¹	
$F(000)$	724	
Crystal color, morphology	colorless, plate	
Crystal size	0.35 x 0.30 x 0.02 mm ³	
Theta range for data collection	0.95 to 25.08°	
Index ranges	$-12 \leq h \leq 12, -9 \leq k \leq 9, -25 \leq l \leq 23$	
Reflections collected	13985	
Independent reflections	3487 [$R(\text{int}) = 0.0746$]	
Observed reflections	2408	
Completeness to theta = 25.08°	99.6%	
Absorption correction	Multi-scan	
Max. and min. transmission	0.9982 and 0.9697	
Refinement method	Full-matrix least-squares on F^2	
Data / restraints / parameters	3487 / 1 / 408	
Goodness-of-fit on F^2	1.016	
Final R indices [$I > 2\sigma(I)$]	$R1 = 0.0527, wR2 = 0.1310$	
R indices (all data)	$R1 = 0.1022, wR2 = 0.1700$	
Absolute structure parameter	0(10)	
Largest diff. peak and hole	0.314 and -0.460 e.Å ⁻³	

Table 2. Atomic coordinates ($\times 10^4$) and equivalent isotropic displacement parameters ($\text{\AA}^2 \times 10^3$) for 08001. U_{eq} is defined as one third of the trace of the orthogonalized U_{ij} tensor.

	x	y	z	U_{eq}
O1	1311(4)	707(5)	7139(2)	29(1)
O2	2682(4)	-818(5)	6198(2)	27(1)
O3	5308(3)	1909(5)	6554(2)	25(1)
O4	4608(4)	1994(5)	5087(2)	26(1)
O5	5152(3)	4703(5)	5785(2)	24(1)
N1	3562(4)	4120(6)	6867(2)	21(1)
C1	2623(6)	3056(7)	7163(3)	25(1)
C2	2174(5)	1622(7)	6764(3)	20(1)
C3	3249(5)	474(7)	6560(3)	23(1)
C4	4322(5)	1164(7)	6169(3)	22(1)
C5	3844(6)	2274(7)	5627(3)	21(1)
C6	3847(5)	4148(7)	5728(3)	22(1)
C7	3102(5)	4796(8)	6270(3)	27(1)
C8	4153(6)	5330(8)	7303(3)	30(1)
C9	3272(6)	6510(7)	7633(3)	31(2)
C10	4040(6)	7555(9)	8102(3)	38(2)
C11	4481(7)	6607(8)	8680(3)	41(2)
C12	5336(7)	7555(10)	9136(3)	41(2)
C13	5735(7)	6591(10)	9719(3)	49(2)
C14	6641(7)	7494(11)	10157(3)	51(2)
C15	7016(8)	6530(11)	10740(4)	58(2)
C16	7977(8)	7383(14)	11162(3)	65(3)
O6	7638(4)	3831(5)	6277(2)	28(1)
O7	8657(4)	5383(5)	5264(2)	27(1)
O8	10348(3)	7959(5)	6390(2)	23(1)
O9	8998(4)	10016(5)	5252(2)	29(1)
O10	8888(4)	10945(5)	6529(2)	30(1)
N2	8263(4)	7804(6)	7073(2)	24(1)
C17	8018(5)	6049(7)	6959(3)	24(1)
C18	7888(5)	5563(7)	6280(3)	22(1)
C19	9029(5)	5854(7)	5881(3)	22(1)

C20	9552(5)	7630(7)	5854(2)	19(1)
C21	8501(5)	8903(7)	5705(3)	23(1)
C22	7942(5)	9882(7)	6240(3)	22(1)
C23	7326(5)	8878(7)	6744(3)	23(1)
C24	8344(6)	8176(8)	7742(3)	32(2)
C25	9493(6)	7443(9)	8084(3)	36(2)
C26	9543(7)	8005(10)	8759(3)	43(2)
C27	10682(7)	7341(10)	9137(3)	46(2)
C28	10719(6)	7865(12)	9809(3)	50(2)
C29	11839(7)	7229(10)	10196(3)	44(2)
C30	11895(6)	7800(11)	10869(3)	45(2)
C31	13007(7)	7106(10)	11260(3)	49(2)
C32	13006(8)	7702(12)	11926(3)	61(2)
O11	7251(3)	2615(5)	4946(2)	26(1)

Table 3. Bond lengths [\AA] and angles [$^\circ$] for 08001.

O(1)-C(2)	1.435(7)	C(10)-C(11)	1.525(9)
O(1)-H(1C)	0.8400	C(10)-H(10A)	0.9900
O(2)-C(3)	1.430(7)	C(10)-H(10B)	0.9900
O(2)-H(2B)	0.8400	C(11)-C(12)	1.516(9)
O(3)-C(4)	1.432(6)	C(11)-H(11A)	0.9900
O(3)-H(3B)	0.8400	C(11)-H(11B)	0.9900
O(4)-C(5)	1.445(6)	C(12)-C(13)	1.530(10)
O(4)-H(4B)	0.8400	C(12)-H(12A)	0.9900
O(5)-C(6)	1.429(7)	C(12)-H(12B)	0.9900
O(5)-H(5B)	0.8400	C(13)-C(14)	1.504(10)
N(1)-C(1)	1.466(7)	C(13)-H(13A)	0.9900
N(1)-C(7)	1.467(7)	C(13)-H(13B)	0.9900
N(1)-C(8)	1.483(7)	C(14)-C(15)	1.524(10)
C(1)-C(2)	1.520(8)	C(14)-H(14A)	0.9900
C(1)-H(1A)	0.9900	C(14)-H(14B)	0.9900
C(1)-H(1B)	0.9900	C(15)-C(16)	1.499(11)
C(2)-C(3)	1.531(8)	C(15)-H(15A)	0.9900
C(2)-H(2A)	1.0000	C(15)-H(15B)	0.9900
C(3)-C(4)	1.525(8)	C(16)-H(16A)	0.9800
C(3)-H(3A)	1.0000	C(16)-H(16B)	0.9800
C(4)-C(5)	1.548(8)	C(16)-H(16C)	0.9800
C(4)-H(4A)	1.0000	O(6)-C(18)	1.439(7)
C(5)-C(6)	1.547(8)	O(6)-H(6B)	0.8400
C(5)-H(5A)	1.0000	O(7)-C(19)	1.426(7)
C(6)-C(7)	1.515(8)	O(7)-H(7C)	0.8400
C(6)-H(6A)	1.0000	O(8)-C(20)	1.424(6)
C(7)-H(7A)	0.9900	O(8)-H(8C)	0.8400
C(7)-H(7B)	0.9900	O(9)-C(21)	1.441(7)
C(8)-C(9)	1.520(8)	O(9)-H(9C)	0.8400
C(8)-H(8A)	0.9900	O(10)-C(22)	1.438(7)
C(8)-H(8B)	0.9900	O(10)-H(10C)	0.8400
C(9)-C(10)	1.529(8)	N(2)-C(24)	1.473(7)
C(9)-H(9A)	0.9900	N(2)-C(23)	1.475(7)
C(9)-H(9B)	0.9900	N(2)-C(17)	1.476(8)

C(17)-C(18)	1.518(8)	C(31)-C(32)	1.514(10)
C(17)-H(17A)	0.9900	C(31)-H(31A)	0.9900
C(17)-H(17B)	0.9900	C(31)-H(31B)	0.9900
C(18)-C(19)	1.502(8)	C(32)-H(32A)	0.9800
C(18)-H(18A)	1.0000	C(32)-H(32B)	0.9800
C(19)-C(20)	1.552(8)	C(32)-H(32C)	0.9800
C(19)-H(19A)	1.0000	O(11)-H(11C)	1.0679
C(20)-C(21)	1.535(8)	O(11)-H(11D)	0.8495
C(20)-H(20A)	1.0000		
C(21)-C(22)	1.532(8)	C(2)-O(1)-H(1C)	109.5
C(21)-H(21A)	1.0000	C(3)-O(2)-H(2B)	109.5
C(22)-C(23)	1.516(8)	C(4)-O(3)-H(3B)	109.5
C(22)-H(22A)	1.0000	C(5)-O(4)-H(4B)	109.5
C(23)-H(23A)	0.9900	C(6)-O(5)-H(5B)	109.5
C(23)-H(23B)	0.9900	C(1)-N(1)-C(7)	113.8(4)
C(24)-C(25)	1.505(8)	C(1)-N(1)-C(8)	112.8(4)
C(24)-H(24A)	0.9900	C(7)-N(1)-C(8)	114.9(5)
C(24)-H(24B)	0.9900	N(1)-C(1)-C(2)	114.0(4)
C(25)-C(26)	1.524(8)	N(1)-C(1)-H(1A)	108.7
C(25)-H(25A)	0.9900	C(2)-C(1)-H(1A)	108.7
C(25)-H(25B)	0.9900	N(1)-C(1)-H(1B)	108.7
C(26)-C(27)	1.515(9)	C(2)-C(1)-H(1B)	108.7
C(26)-H(26A)	0.9900	H(1A)-C(1)-H(1B)	107.6
C(26)-H(26B)	0.9900	O(1)-C(2)-C(1)	105.6(4)
C(27)-C(28)	1.509(9)	O(1)-C(2)-C(3)	108.3(5)
C(27)-H(27A)	0.9900	C(1)-C(2)-C(3)	115.0(4)
C(27)-H(27B)	0.9900	O(1)-C(2)-H(2A)	109.3
C(28)-C(29)	1.502(9)	C(1)-C(2)-H(2A)	109.3
C(28)-H(28A)	0.9900	C(3)-C(2)-H(2A)	109.3
C(28)-H(28B)	0.9900	O(2)-C(3)-C(4)	105.4(4)
C(29)-C(30)	1.523(9)	O(2)-C(3)-C(2)	108.6(4)
C(29)-H(29A)	0.9900	C(4)-C(3)-C(2)	118.8(5)
C(29)-H(29B)	0.9900	O(2)-C(3)-H(3A)	107.9
C(30)-C(31)	1.517(9)	C(4)-C(3)-H(3A)	107.9
C(30)-H(30A)	0.9900	C(2)-C(3)-H(3A)	107.9
C(30)-H(30B)	0.9900	O(3)-C(4)-C(3)	110.9(4)

O(3)-C(4)-C(5)	113.2(4)	C(11)-C(10)-H(10A)	109.0
C(3)-C(4)-C(5)	114.3(4)	C(9)-C(10)-H(10A)	109.0
O(3)-C(4)-H(4A)	105.9	C(11)-C(10)-H(10B)	109.0
C(3)-C(4)-H(4A)	105.9	C(9)-C(10)-H(10B)	109.0
C(5)-C(4)-H(4A)	105.9	H(10A)-C(10)-H(10B)	107.8
O(4)-C(5)-C(6)	105.8(4)	C(12)-C(11)-C(10)	115.1(6)
O(4)-C(5)-C(4)	110.1(4)	C(12)-C(11)-H(11A)	108.5
C(6)-C(5)-C(4)	118.4(5)	C(10)-C(11)-H(11A)	108.5
O(4)-C(5)-H(5A)	107.4	C(12)-C(11)-H(11B)	108.5
C(6)-C(5)-H(5A)	107.4	C(10)-C(11)-H(11B)	108.5
C(4)-C(5)-H(5A)	107.4	H(11A)-C(11)-H(11B)	107.5
O(5)-C(6)-C(7)	109.0(5)	C(11)-C(12)-C(13)	114.0(6)
O(5)-C(6)-C(5)	108.9(5)	C(11)-C(12)-H(12A)	108.7
C(7)-C(6)-C(5)	117.1(5)	C(13)-C(12)-H(12A)	108.7
O(5)-C(6)-H(6A)	107.1	C(11)-C(12)-H(12B)	108.7
C(7)-C(6)-H(6A)	107.1	C(13)-C(12)-H(12B)	108.7
C(5)-C(6)-H(6A)	107.1	H(12A)-C(12)-H(12B)	107.6
N(1)-C(7)-C(6)	112.7(5)	C(14)-C(13)-C(12)	114.2(7)
N(1)-C(7)-H(7A)	109.1	C(14)-C(13)-H(13A)	108.7
C(6)-C(7)-H(7A)	109.1	C(12)-C(13)-H(13A)	108.7
N(1)-C(7)-H(7B)	109.1	C(14)-C(13)-H(13B)	108.7
C(6)-C(7)-H(7B)	109.1	C(12)-C(13)-H(13B)	108.7
H(7A)-C(7)-H(7B)	107.8	H(13A)-C(13)-H(13B)	107.6
N(1)-C(8)-C(9)	118.5(5)	C(13)-C(14)-C(15)	113.7(7)
N(1)-C(8)-H(8A)	107.7	C(13)-C(14)-H(14A)	108.8
C(9)-C(8)-H(8A)	107.7	C(15)-C(14)-H(14A)	108.8
N(1)-C(8)-H(8B)	107.7	C(13)-C(14)-H(14B)	108.8
C(9)-C(8)-H(8B)	107.7	C(15)-C(14)-H(14B)	108.8
H(8A)-C(8)-H(8B)	107.1	H(14A)-C(14)-H(14B)	107.7
C(8)-C(9)-C(10)	110.9(5)	C(16)-C(15)-C(14)	114.0(8)
C(8)-C(9)-H(9A)	109.5	C(16)-C(15)-H(15A)	108.7
C(10)-C(9)-H(9A)	109.5	C(14)-C(15)-H(15A)	108.7
C(8)-C(9)-H(9B)	109.5	C(16)-C(15)-H(15B)	108.7
C(10)-C(9)-H(9B)	109.5	C(14)-C(15)-H(15B)	108.7
H(9A)-C(9)-H(9B)	108.0	H(15A)-C(15)-H(15B)	107.6
C(11)-C(10)-C(9)	113.1(6)	C(15)-C(16)-H(16A)	109.5

C(15)-C(16)-H(16B)	109.5	C(19)-C(20)-H(20A)	105.7
H(16A)-C(16)-H(16B)	109.5	O(9)-C(21)-C(22)	109.4(4)
C(15)-C(16)-H(16C)	109.5	O(9)-C(21)-C(20)	107.7(4)
H(16A)-C(16)-H(16C)	109.5	C(22)-C(21)-C(20)	118.6(5)
H(16B)-C(16)-H(16C)	109.5	O(9)-C(21)-H(21A)	106.9
C(18)-O(6)-H(6B)	109.5	C(22)-C(21)-H(21A)	106.9
C(19)-O(7)-H(7C)	109.5	C(20)-C(21)-H(21A)	106.9
C(20)-O(8)-H(8C)	109.5	O(10)-C(22)-C(23)	108.3(5)
C(21)-O(9)-H(9C)	109.5	O(10)-C(22)-C(21)	111.9(4)
C(22)-O(10)-H(10C)	109.5	C(23)-C(22)-C(21)	115.7(5)
C(24)-N(2)-C(23)	111.4(4)	O(10)-C(22)-H(22A)	106.8
C(24)-N(2)-C(17)	111.6(5)	C(23)-C(22)-H(22A)	106.8
C(23)-N(2)-C(17)	113.0(5)	C(21)-C(22)-H(22A)	106.8
N(2)-C(17)-C(18)	115.1(5)	N(2)-C(23)-C(22)	112.4(5)
N(2)-C(17)-H(17A)	108.5	N(2)-C(23)-H(23A)	109.1
C(18)-C(17)-H(17A)	108.5	C(22)-C(23)-H(23A)	109.1
N(2)-C(17)-H(17B)	108.5	N(2)-C(23)-H(23B)	109.1
C(18)-C(17)-H(17B)	108.5	C(22)-C(23)-H(23B)	109.1
H(17A)-C(17)-H(17B)	107.5	H(23A)-C(23)-H(23B)	107.9
O(6)-C(18)-C(19)	107.3(5)	N(2)-C(24)-C(25)	114.7(5)
O(6)-C(18)-C(17)	105.9(5)	N(2)-C(24)-H(24A)	108.6
C(19)-C(18)-C(17)	117.5(5)	C(25)-C(24)-H(24A)	108.6
O(6)-C(18)-H(18A)	108.6	N(2)-C(24)-H(24B)	108.6
C(19)-C(18)-H(18A)	108.6	C(25)-C(24)-H(24B)	108.6
C(17)-C(18)-H(18A)	108.6	H(24A)-C(24)-H(24B)	107.6
O(7)-C(19)-C(18)	107.2(4)	C(24)-C(25)-C(26)	110.5(5)
O(7)-C(19)-C(20)	107.6(4)	C(24)-C(25)-H(25A)	109.6
C(18)-C(19)-C(20)	116.9(5)	C(26)-C(25)-H(25A)	109.6
O(7)-C(19)-H(19A)	108.3	C(24)-C(25)-H(25B)	109.6
C(18)-C(19)-H(19A)	108.3	C(26)-C(25)-H(25B)	109.6
C(20)-C(19)-H(19A)	108.3	H(25A)-C(25)-H(25B)	108.1
O(8)-C(20)-C(21)	115.6(4)	C(27)-C(26)-C(25)	114.1(6)
O(8)-C(20)-C(19)	109.9(4)	C(27)-C(26)-H(26A)	108.7
C(21)-C(20)-C(19)	113.3(4)	C(25)-C(26)-H(26A)	108.7
O(8)-C(20)-H(20A)	105.7	C(27)-C(26)-H(26B)	108.7
C(21)-C(20)-H(20A)	105.7	C(25)-C(26)-H(26B)	108.7

H(26A)-C(26)-H(26B)	107.6	C(31)-C(30)-H(30A)	108.6
C(28)-C(27)-C(26)	114.3(6)	C(29)-C(30)-H(30A)	108.6
C(28)-C(27)-H(27A)	108.7	C(31)-C(30)-H(30B)	108.6
C(26)-C(27)-H(27A)	108.7	C(29)-C(30)-H(30B)	108.6
C(28)-C(27)-H(27B)	108.7	H(30A)-C(30)-H(30B)	107.5
C(26)-C(27)-H(27B)	108.7	C(32)-C(31)-C(30)	112.6(7)
H(27A)-C(27)-H(27B)	107.6	C(32)-C(31)-H(31A)	109.1
C(29)-C(28)-C(27)	115.5(6)	C(30)-C(31)-H(31A)	109.1
C(29)-C(28)-H(28A)	108.4	C(32)-C(31)-H(31B)	109.1
C(27)-C(28)-H(28A)	108.4	C(30)-C(31)-H(31B)	109.1
C(29)-C(28)-H(28B)	108.4	H(31A)-C(31)-H(31B)	107.8
C(27)-C(28)-H(28B)	108.4	C(31)-C(32)-H(32A)	109.5
H(28A)-C(28)-H(28B)	107.5	C(31)-C(32)-H(32B)	109.5
C(28)-C(29)-C(30)	115.4(6)	H(32A)-C(32)-H(32B)	109.5
C(28)-C(29)-H(29A)	108.4	C(31)-C(32)-H(32C)	109.5
C(30)-C(29)-H(29A)	108.4	H(32A)-C(32)-H(32C)	109.5
C(28)-C(29)-H(29B)	108.4	H(32B)-C(32)-H(32C)	109.5
C(30)-C(29)-H(29B)	108.4	H(11C)-O(11)-H(11D)	102.2
H(29A)-C(29)-H(29B)	107.5		
C(31)-C(30)-C(29)	114.9(6)		

Symmetry transformations used to generate equivalent atoms:

Table 4. Anisotropic displacement parameters ($\text{\AA}^2 \times 10^3$) for 08001. The anisotropic displacement factor exponent takes the form: $-2\pi^2 [h^2 a^{*2} U_{11} + \dots + 2 h k a^* b^* U_{12}]$

	U_{11}	U_{22}	U_{33}	U_{23}	U_{13}	U_{12}
O1	35(2)	22(2)	31(2)	-3(2)	8(2)	-8(2)
O2	26(2)	22(2)	34(2)	-3(2)	5(2)	-3(2)
O3	20(2)	22(2)	33(2)	5(2)	-6(2)	-4(2)
O4	26(2)	28(2)	26(2)	-6(2)	8(2)	-5(2)
O5	21(2)	20(2)	30(2)	3(2)	-5(2)	-1(2)
N1	27(3)	15(2)	21(2)	-6(2)	4(2)	-5(2)
C1	30(3)	20(3)	25(3)	-8(3)	5(3)	-6(3)
C2	15(3)	22(3)	24(3)	-6(2)	4(2)	-5(2)
C3	28(3)	18(3)	21(3)	-4(2)	-3(2)	-4(2)
C4	18(3)	21(3)	28(3)	-1(3)	4(2)	-3(2)
C5	26(3)	19(3)	19(3)	0(2)	7(2)	0(2)
C6	15(3)	23(3)	28(3)	4(3)	-2(2)	0(2)
C7	27(3)	19(3)	34(4)	-6(3)	2(3)	0(3)
C8	29(3)	27(3)	33(3)	-12(3)	-2(3)	-4(3)
C9	31(3)	23(3)	39(4)	-6(3)	-1(3)	3(3)
C10	46(4)	27(3)	40(4)	-9(3)	-3(3)	-3(3)
C11	51(4)	28(4)	43(4)	-6(3)	-5(3)	-2(3)
C12	46(4)	45(4)	32(4)	-10(4)	2(3)	-4(4)
C13	56(5)	42(4)	49(4)	-10(4)	-14(4)	7(4)
C14	51(4)	62(5)	39(4)	-3(4)	-3(3)	-6(4)
C15	61(5)	62(6)	49(5)	-7(4)	-16(4)	10(4)
C16	57(5)	103(8)	35(4)	-3(5)	2(4)	-8(5)
O6	19(2)	14(2)	49(3)	2(2)	-3(2)	-4(2)
O7	26(2)	22(2)	33(2)	-8(2)	0(2)	-6(2)
O8	21(2)	22(2)	27(2)	-1(2)	-3(2)	-4(2)
O9	24(2)	27(2)	37(3)	11(2)	8(2)	1(2)
O10	27(2)	18(2)	43(3)	-3(2)	-6(2)	-4(2)
N2	29(3)	21(3)	22(3)	-3(2)	0(2)	2(2)
C17	21(3)	20(3)	30(3)	6(3)	2(2)	1(2)
C18	22(3)	12(3)	32(3)	1(3)	0(3)	-3(2)
C19	20(3)	19(3)	28(3)	-4(3)	0(2)	-5(2)

C20	21(3)	11(3)	25(3)	-1(3)	0(2)	-1(2)
C21	24(3)	17(3)	26(3)	3(3)	3(2)	-2(3)
C22	20(3)	15(3)	29(3)	2(3)	-3(3)	0(2)
C23	24(3)	24(3)	21(3)	-1(3)	0(2)	6(3)
C24	35(3)	38(4)	23(3)	-1(3)	4(3)	7(3)
C25	40(4)	39(4)	29(3)	-5(3)	-3(3)	8(3)
C26	47(4)	57(5)	25(3)	-6(4)	-3(3)	3(4)
C27	49(4)	50(5)	37(4)	1(3)	-7(3)	8(4)
C28	39(4)	76(6)	35(4)	-5(4)	-5(3)	5(4)
C29	45(4)	55(5)	33(4)	-2(3)	-10(3)	3(4)
C30	48(4)	56(5)	32(4)	-1(4)	-3(3)	3(4)
C31	60(5)	50(5)	35(4)	-2(4)	-12(4)	-6(4)
C32	87(6)	57(5)	37(4)	-5(4)	-21(4)	-12(5)
O11	29(2)	19(2)	30(2)	0(2)	2(2)	0(2)

Table 5. Hydrogen coordinates ($\times 10^4$) and isotropic displacement parameters ($\text{\AA}^2 \times 10^3$) for 08001.

	x	y	z	U(eq)
H1C	1209	-234	6989	44
H2B	1947	-1036	6329	41
H3B	5470	2847	6418	37
H4B	5374	2282	5164	39
H5B	5578	4068	6019	36
H1A	1861	3718	7270	30
H1B	3014	2628	7556	30
H2A	1691	2043	6388	24
H3A	3658	-22	6942	27
H4A	4737	199	5972	27
H5A	2937	1943	5516	25
H6A	3460	4654	5343	26
H7A	3181	6002	6283	32
H7B	2177	4525	6205	32
H8A	4776	5987	7069	36
H8B	4655	4718	7625	36
H9A	2603	5886	7850	38
H9B	2832	7229	7323	38
H10A	4807	8006	7900	45
H10B	3498	8487	8229	45
H11A	3707	6232	8898	49
H11B	4955	5622	8547	49
H12A	4873	8555	9262	49
H12B	6124	7902	8924	49
H13A	4948	6297	9944	59
H13B	6153	5561	9592	59
H14A	6229	8532	10281	61
H14B	7434	7773	9935	61
H15A	6227	6305	10974	69
H15B	7381	5465	10616	69

H16A	8256	6640	11497	98
H16B	7576	8354	11340	98
H16C	8726	7711	10925	98
H6B	6884	3658	6139	41
H7C	8242	4501	5274	41
H8C	9982	7617	6708	35
H9C	9777	9797	5193	44
H10C	9350	11355	6256	44
H17A	7215	5739	7167	28
H17B	8733	5413	7156	28
H18A	7122	6137	6090	27
H19A	9747	5123	6030	26
H20A	10142	7655	5496	23
H21A	7764	8306	5498	27
H22A	7250	10599	6055	26
H23A	6934	9625	7046	28
H23B	6627	8201	6554	28
H24A	7549	7776	7935	38
H24B	8372	9378	7795	38
H25A	10294	7778	7880	43
H25B	9436	6235	8068	43
H26A	9576	9215	8768	52
H26B	8737	7662	8956	52
H27A	10660	6132	9119	55
H27B	11488	7705	8944	55
H28A	10735	9074	9824	60
H28B	9910	7501	9999	60
H29A	12648	7564	9999	53
H29B	11808	6019	10191	53
H30A	11957	9008	10874	54
H30B	11075	7497	11063	54
H31A	12952	5897	11257	59
H31B	13831	7421	11073	59
H32A	13740	7222	12158	91
H32B	12200	7373	12116	91
H32C	13079	8897	11933	91

H11C	7564	2984	4500	39
H11D	7769	1840	5037	39

Table 6. Torsion angles [°] for 08001.

C7-N1-C1-C2	56.9(6)	C23-N2-C17-C18	52.1(6)
C8-N1-C1-C2	-169.9(5)	N2-C17-C18-O6	-179.2(4)
N1-C1-C2-O1	177.9(5)	N2-C17-C18-C19	61.0(7)
N1-C1-C2-C3	58.6(7)	O6-C18-C19-O7	63.8(6)
O1-C2-C3-O2	61.4(5)	C17-C18-C19-O7	-177.1(5)
C1-C2-C3-O2	179.2(5)	O6-C18-C19-C20	-175.3(5)
O1-C2-C3-C4	-178.2(4)	C17-C18-C19-C20	-56.3(7)
C1-C2-C3-C4	-60.5(7)	O7-C19-C20-O8	-159.1(4)
O2-C3-C4-O3	-154.1(4)	C18-C19-C20-O8	80.2(6)
C2-C3-C4-O3	84.0(6)	O7-C19-C20-C21	69.8(5)
O2-C3-C4-C5	76.5(6)	C18-C19-C20-C21	-50.8(7)
C2-C3-C4-C5	-45.5(7)	O8-C20-C21-O9	95.9(5)
O3-C4-C5-O4	90.0(5)	C19-C20-C21-O9	-135.9(5)
C3-C4-C5-O4	-141.7(5)	O8-C20-C21-C22	-28.9(7)
O3-C4-C5-C6	-31.8(7)	C19-C20-C21-C22	99.3(6)
C3-C4-C5-C6	96.5(6)	O9-C21-C22-O10	-57.8(6)
O4-C5-C6-O5	-56.0(6)	C20-C21-C22-O10	66.1(6)
C4-C5-C6-O5	67.9(6)	O9-C21-C22-C23	177.5(4)
O4-C5-C6-C7	179.7(4)	C20-C21-C22-C23	-58.6(7)
C4-C5-C6-C7	-56.3(7)	C24-N2-C23-C22	120.4(5)
C1-N1-C7-C6	-113.2(5)	C17-N2-C23-C22	-113.0(5)
C8-N1-C7-C6	114.6(5)	O10-C22-C23-N2	-63.0(6)
O5-C6-C7-N1	-65.5(6)	C21-C22-C23-N2	63.5(7)
C5-C6-C7-N1	58.7(7)	C23-N2-C24-C25	-165.6(5)
C1-N1-C8-C9	-58.1(7)	C17-N2-C24-C25	67.0(7)
C7-N1-C8-C9	74.5(7)	N2-C24-C25-C26	175.7(6)
N1-C8-C9-C10	175.0(5)	C24-C25-C26-C27	-179.2(6)
C8-C9-C10-C11	-73.0(7)	C25-C26-C27-C28	-178.7(7)
C9-C10-C11-C12	175.3(6)	C26-C27-C28-C29	-179.9(7)
C10-C11-C12-C13	178.4(6)	C27-C28-C29-C30	178.3(7)
C11-C12-C13-C14	176.9(6)	C28-C29-C30-C31	178.1(7)
C12-C13-C14-C15	179.2(6)	C29-C30-C31-C32	-179.5(7)
C13-C14-C15-C16	176.8(7)		
C24-N2-C17-C18	178.6(4)		

Table 7. Hydrogen bonds for 08001 [\AA and $^\circ$].

D-H...A	d(D-H)	d(H...A)	d(D...A)	$\angle(\text{DHA})$
O1-H1C...O8#1	0.84	2.14	2.924(5)	155.5
O1-H1C...O2	0.84	2.37	2.805(5)	112.7
O2-H2B...O8#1	0.84	1.86	2.662(5)	159.6
O2-H2B...O1	0.84	2.36	2.805(5)	113.5
O3-H3B...O5	0.84	2.06	2.825(5)	150.9
O3-H3B...O6	0.84	2.41	2.958(5)	123.0
O4-H4B...O11	0.84	2.03	2.811(5)	153.8
O5-H5B...O3	0.84	2.13	2.825(5)	139.6
O5-H5B...O6	0.84	2.20	2.847(5)	134.0
O6-H6B...O5	0.84	2.11	2.847(5)	146.3
O6-H6B...O3	0.84	2.37	2.958(5)	127.8
O7-H7C...O11	0.84	1.97	2.767(5)	157.6
O7-H7C...O6	0.84	2.33	2.762(6)	112.1
O8-H8C...N2	0.84	1.98	2.654(6)	137.2
O9-H9C...O7#2	0.84	1.98	2.719(5)	145.9
O10-H10C...O9	0.84	2.44	2.861(6)	111.7
O11-H11C...O2#3	1.07	1.81	2.781(5)	149.8
O11-H11D...O9#4	0.85	2.01	2.856(6)	179.8

Symmetry transformations used to generate equivalent atoms:

#1 $x-1, y-1, z$ #2 $-x+2, y+1/2, -z+1$ #3 $-x+1, y+1/2, -z+1$

#4 $x, y-1, z$

checkCIF/PLATON report

No syntax errors found. CIF dictionary Interpreting this report

Datablock: 08001

Bond precision: C-C = 0.0092 A Wavelength=0.71073

Cell: a=10.3632(13) b=8.1773(10) c=21.547(3)
 alpha=90 beta=91.751(2) gamma=90

	Calculated	Reported
Volume	1825.1(4)	1825.1(4)
Space group	P 21	P2(1)
Hall group	P 2yb	?
Moiety formula	2(C ₁₆ H ₃₃ NO ₅), H ₂ O	2(C ₁₆ H ₃₃ NO ₅), H ₂ O
Sum formula	C ₃₂ H ₆₈ N ₂ O ₁₁	C ₃₂ H ₆₈ N ₂ O ₁₁
Mr	656.88	656.88
Dx,g cm ⁻³	1.195	1.195
Z	2	2
Mu (mm ⁻¹)	0.088	0.088
F000	724.0	724.0
F000'	724.38	
h,k,lmax	12,9,25	12,9,25
Nref	3501(6500)	3487
Tmin,Tmax	0.970,0.998	0.970,0.998
Tmin'	0.970	

Correction method= AbsCorr=MULTI-SCAN

Data completeness= 1.00(0.54) Theta(max)= 25.080

R(reflections)= 0.0527(2408) wR2(reflections)= 0.1700(3487)

S = 1.016 Npar= 408

The following ALERTS were generated. Each ALERT has the format test-name_ALERT_alert-type_alert-level.

Alert level B

PLAT222_ALERT_3_B Large Non-Solvent H Ueq(max)/Ueq(min) ... 4.08 Ratio
PLAT416_ALERT_2_B Short Intra D-H..H-D H1C .. H2B .. 1.76 Ang.
PLAT416_ALERT_2_B Short Intra D-H..H-D H3B .. H5B .. 1.32 Ang.
PLAT417_ALERT_2_B Short Inter D-H..H-D H2B .. H11C .. 2.04 Ang.
PLAT417_ALERT_2_B Short Inter D-H..H-D H3B .. H6B .. 1.73 Ang.
PLAT417_ALERT_2_B Short Inter D-H..H-D H5B .. H6B .. 1.41 Ang.

Alert level C

STRVA01_ALERT_4_C Flack test results are meaningless.
From the CIF: `_refine_ls_abs_structure_Flack` 0.000
From the CIF: `_refine_ls_abs_structure_Flack_su` 10.000
PLAT032_ALERT_4_C Std. Uncertainty in Flack Parameter too High ... 10.00
PLAT066_ALERT_1_C Predicted and Reported Transmissions Identical . ?
PLAT220_ALERT_2_C Large Non-Solvent C Ueq(max)/Ueq(min) ... 3.20 Ratio
PLAT220_ALERT_2_C Large Non-Solvent C Ueq(max)/Ueq(min) ... 3.21 Ratio
PLAT222_ALERT_3_C Large Non-Solvent H Ueq(max)/Ueq(min) ... 3.96 Ratio
PLAT340_ALERT_3_C Low Bond Precision on C-C Bonds (x 1000) Ang ... 9
PLAT355_ALERT_3_C Long O-H Bond (0.82A) O11 - H11C ... 1.07 Ang.
PLAT414_ALERT_2_C Short Intra D-H..H-X H9C .. H20A .. 1.90 Ang.
PLAT850_ALERT_2_C Check Flack Parameter Exact Value 0.00 and su .. 10.00

Alert level G

REFLT03_ALERT_4_G Please check that the estimate of the number of Friedel pairs is correct. If it is not, please give the correct count in the `_publ_section_exptl_refinement` section of the submitted CIF.

From the CIF: `_diffn_refl_theta_max` 25.08
From the CIF: `_reflns_number_total` 3487
Count of symmetry unique reflns 3501
Completeness (`_total/calc`) 99.60%
TEST3: Check Friedels for noncentro structure
Estimate of Friedel pairs measured 0
Fraction of Friedel pairs measured 0.000
Are heavy atom types Z>Si present no
PLAT791_ALERT_1_G Confirm the Absolute Configuration of C2 ... S
PLAT791_ALERT_1_G Confirm the Absolute Configuration of C3 ... R
PLAT791_ALERT_1_G Confirm the Absolute Configuration of C5 ... R
PLAT791_ALERT_1_G Confirm the Absolute Configuration of C6 ... R
PLAT791_ALERT_1_G Confirm the Absolute Configuration of C18 ... S
PLAT791_ALERT_1_G Confirm the Absolute Configuration of C19 ... R
PLAT791_ALERT_1_G Confirm the Absolute Configuration of C21 ... R
PLAT791_ALERT_1_G Confirm the Absolute Configuration of C22 ... R

PLAT860_ALERT_3_G Note: Number of Least-Squares Restraints 1

0 ALERT level A = In general: serious problem

6 ALERT level B = Potentially serious problem

10 ALERT level C = Check and explain

10 ALERT level G = General alerts; check

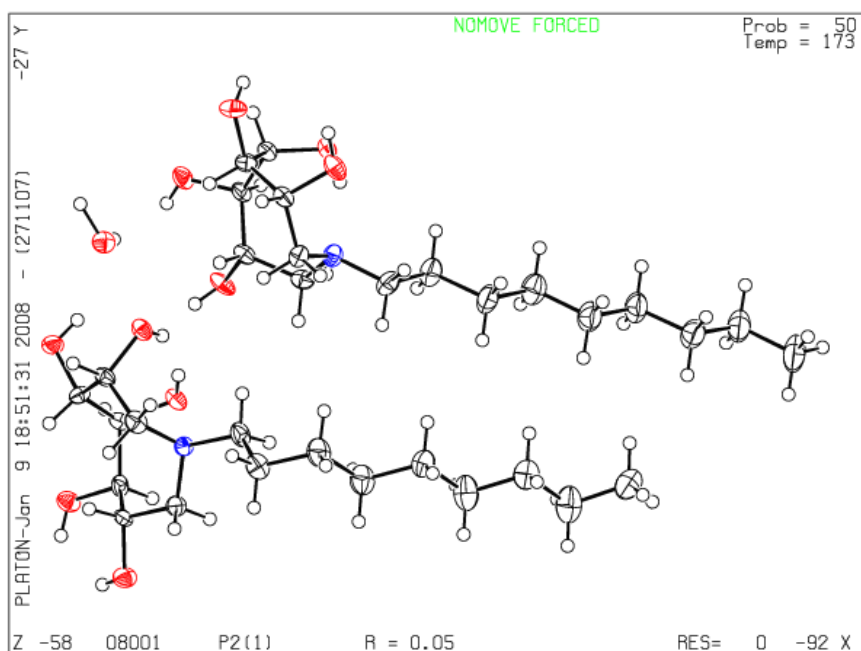
9 ALERT type 1 CIF construction/syntax error, inconsistent or missing data

9 ALERT type 2 Indicator that the structure model may be wrong or deficient

5 ALERT type 3 Indicator that the structure quality may be low

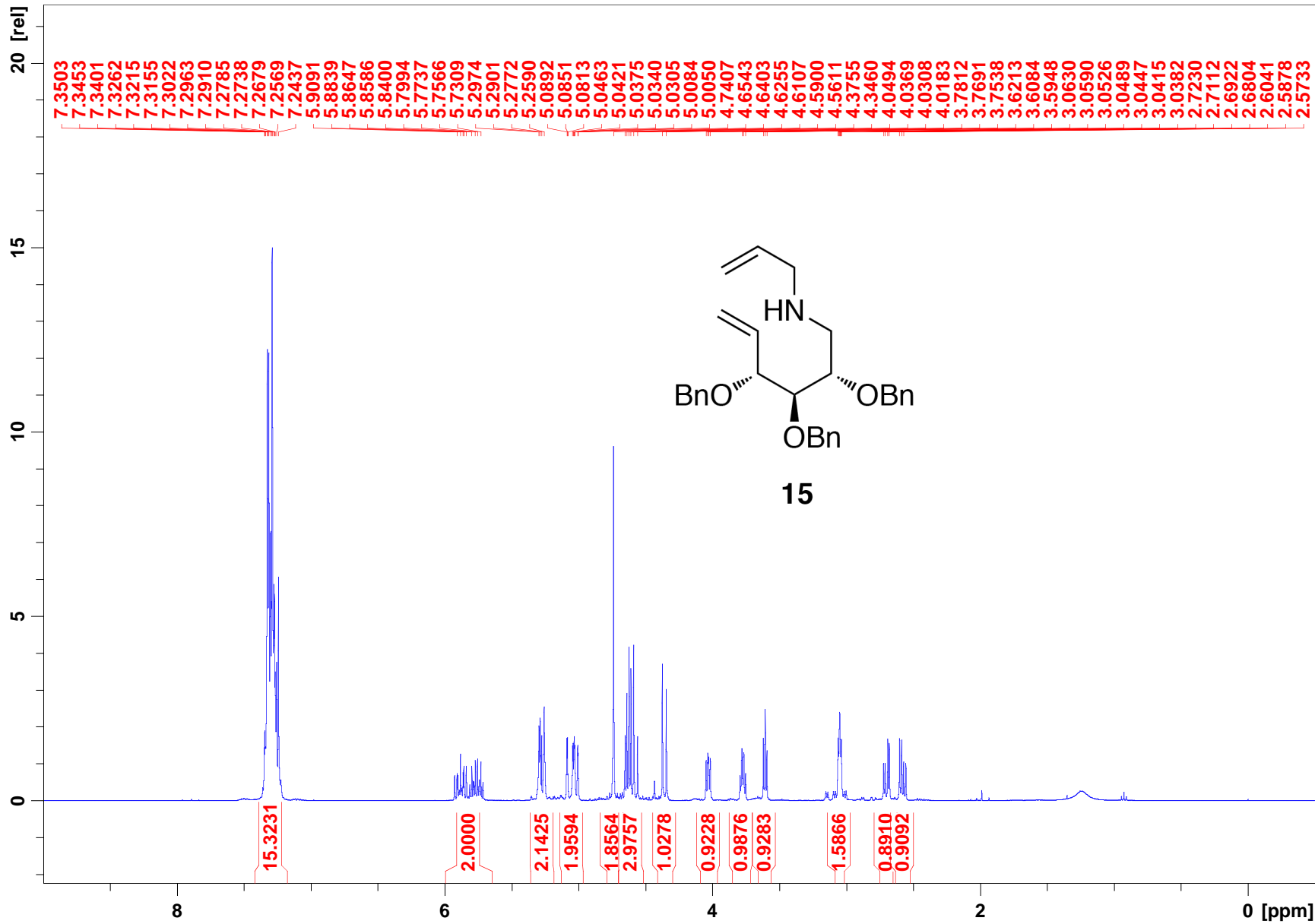
3 ALERT type 4 Improvement, methodology, query or suggestion

0 ALERT type 5 Informative message, check



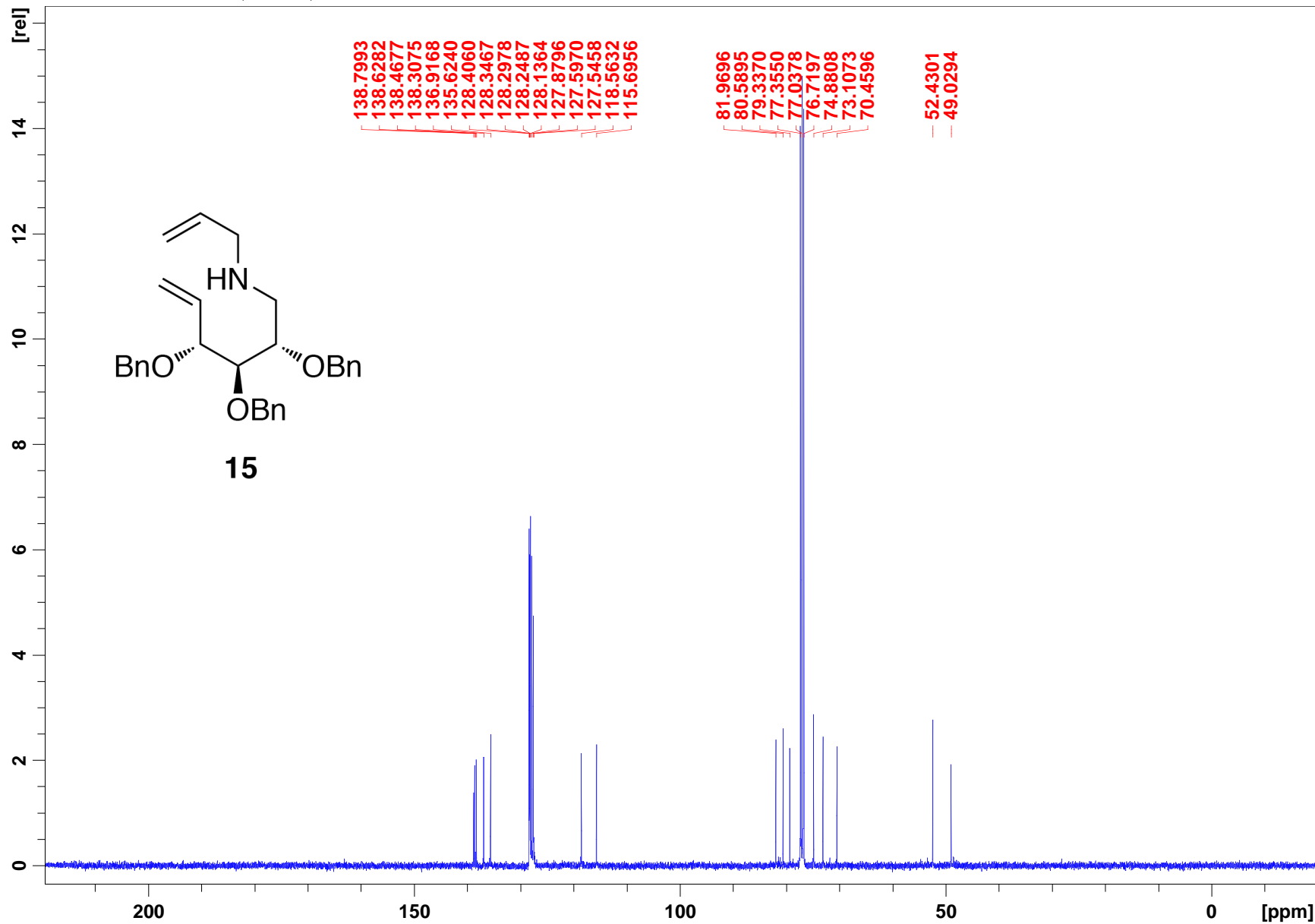
1H-NMR

JL-090307 1 1 C:\Bruker\TOPSPIN leex3451



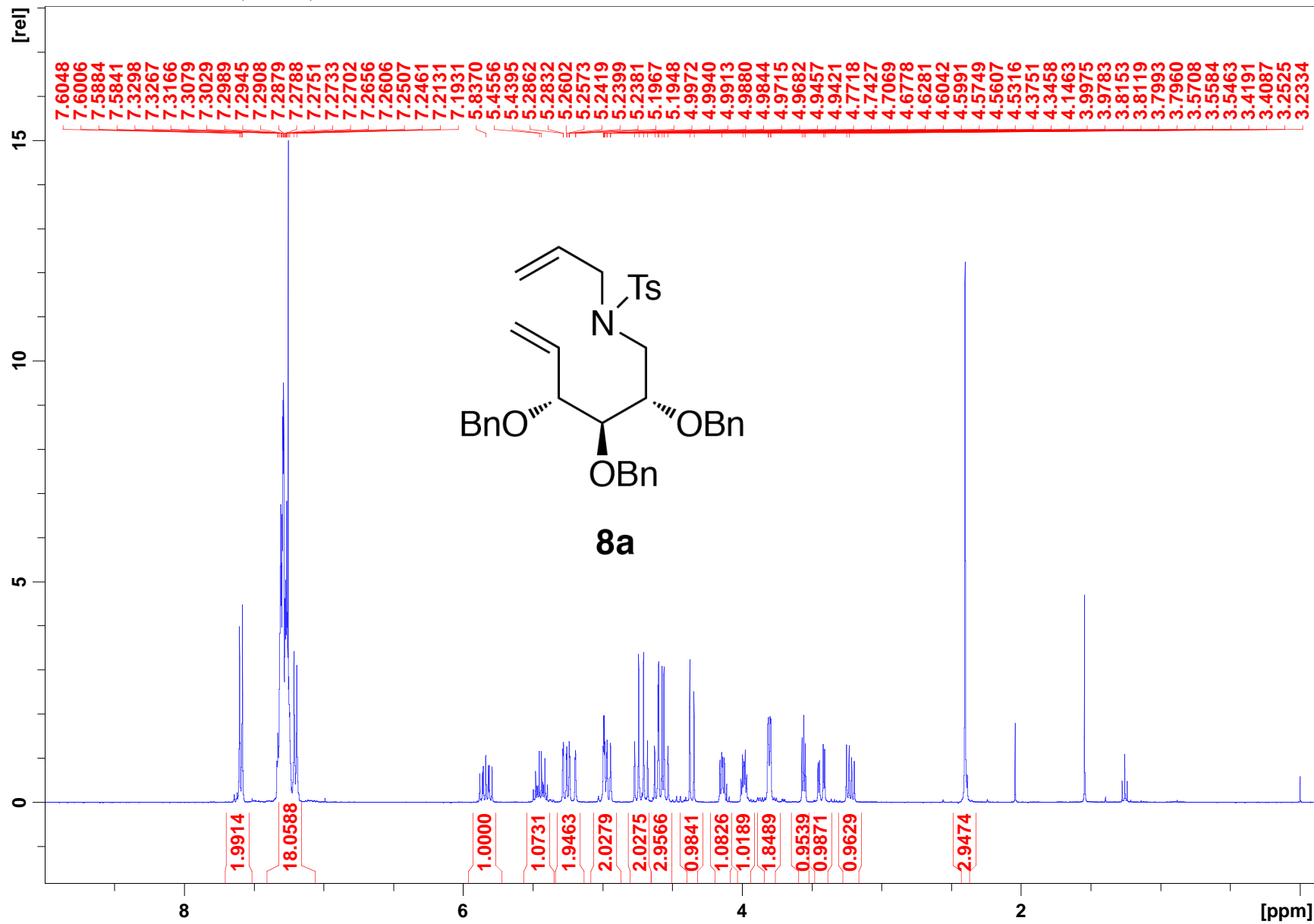
13C-NMR

JL-090307 3 1 C:\Bruker\TOPSPIN leex3451



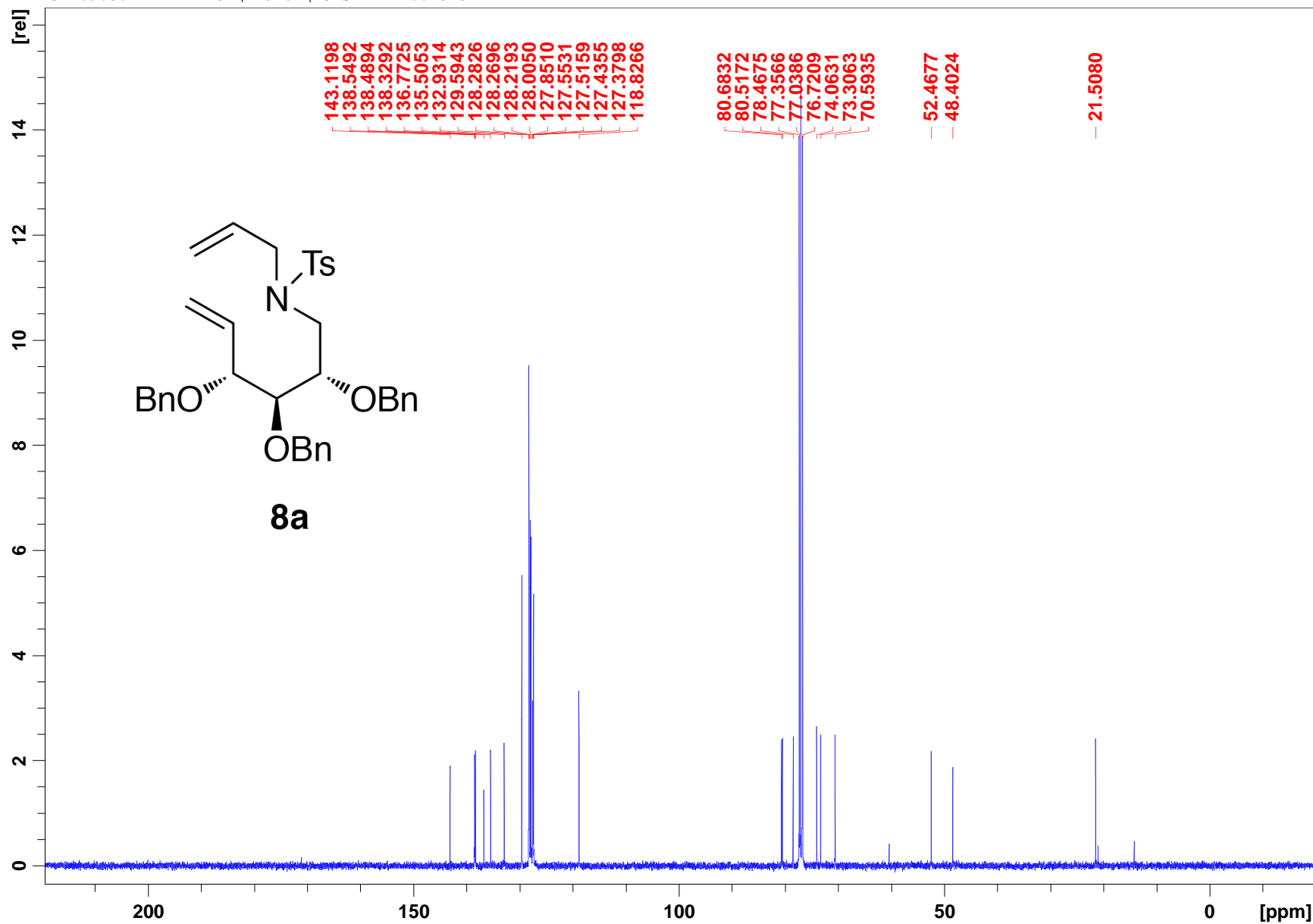
1H-NMR

JL-090507 1 1 C:\Bruker\TOPSPIN leex3451



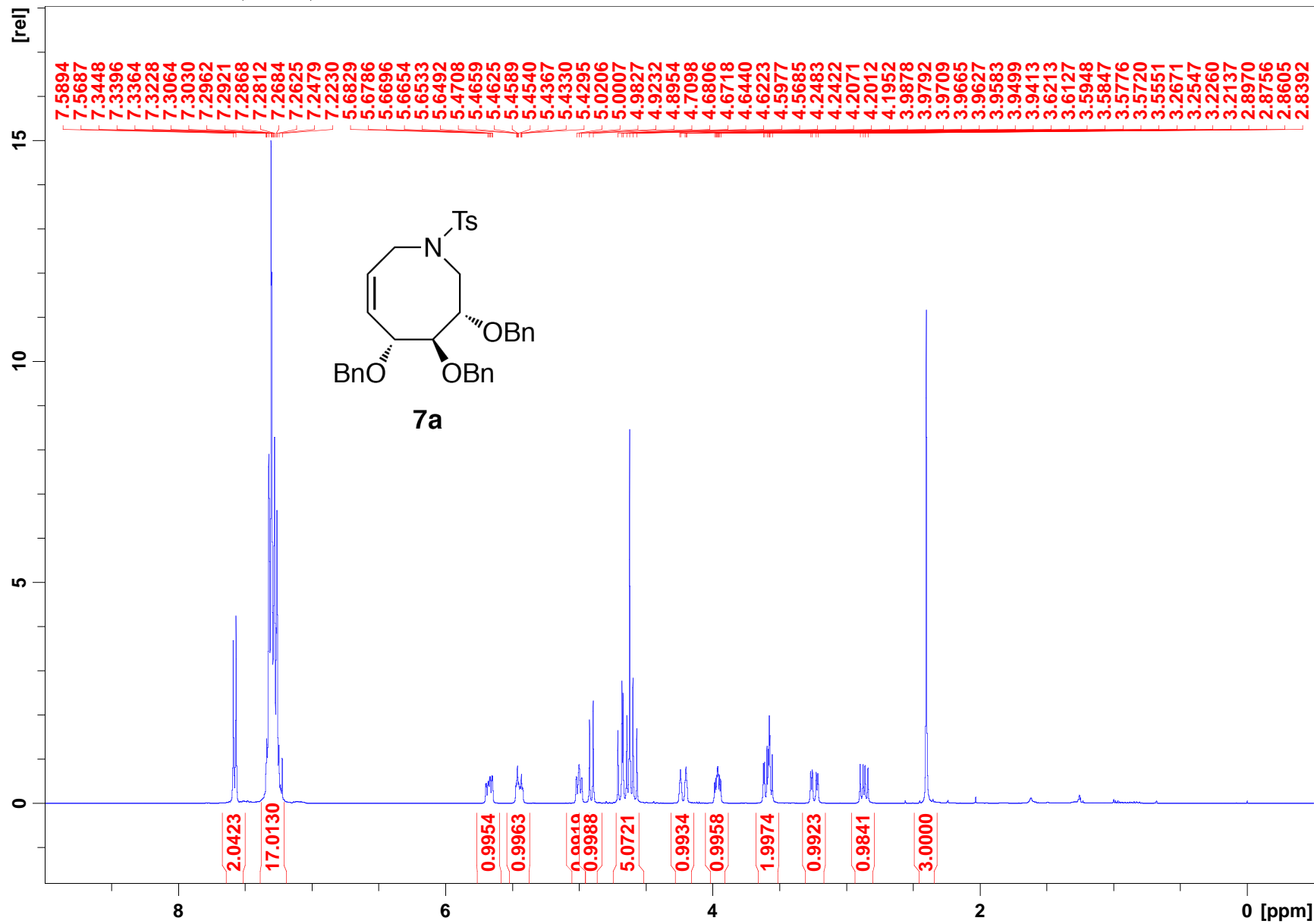
13C-NMR

JL-090507 4 1 C:\Bruker\TOPSPIN leex3451



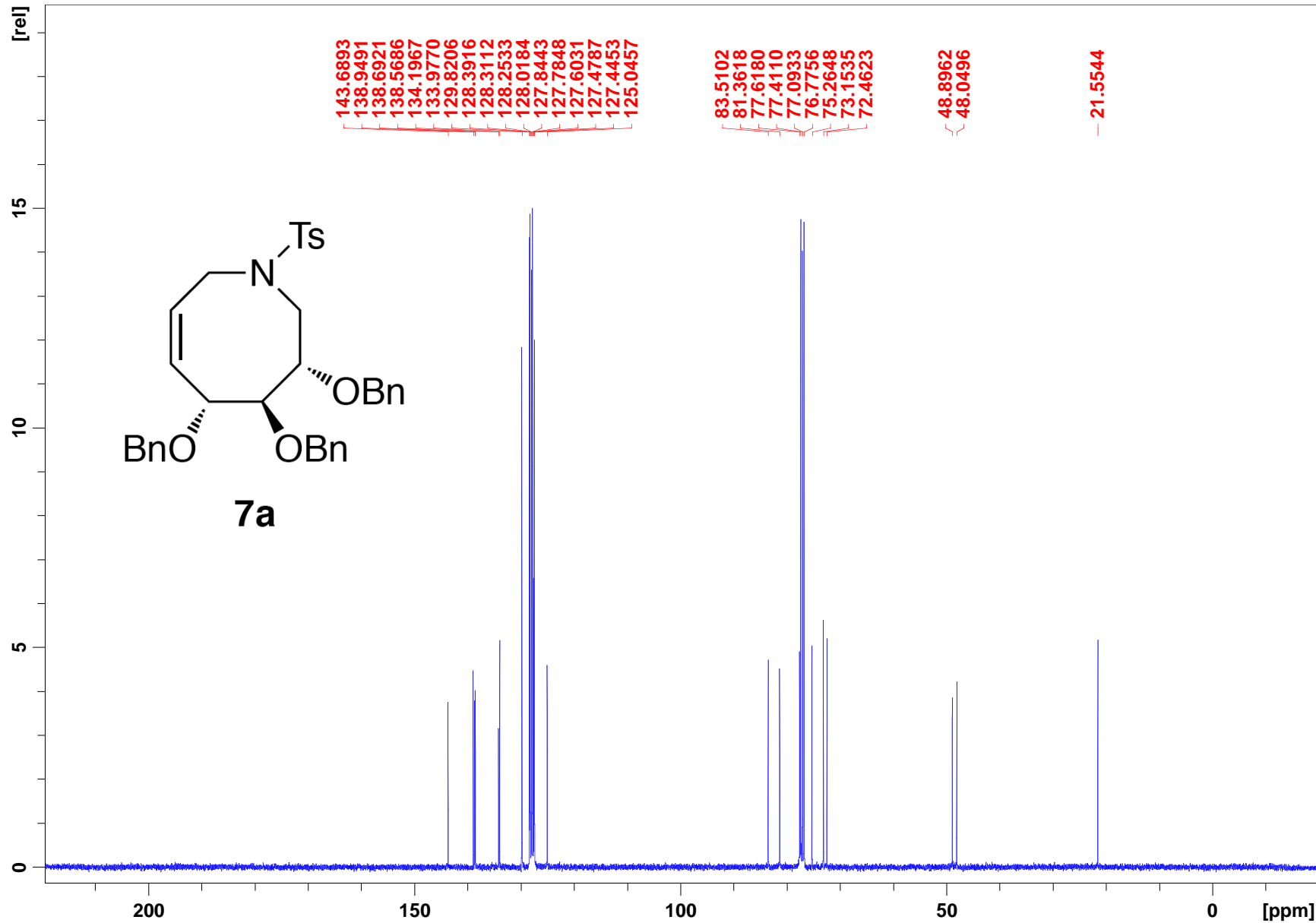
1H-NMR

JL-062207 1 1 C:\Bruker\TOPSPIN leex3451



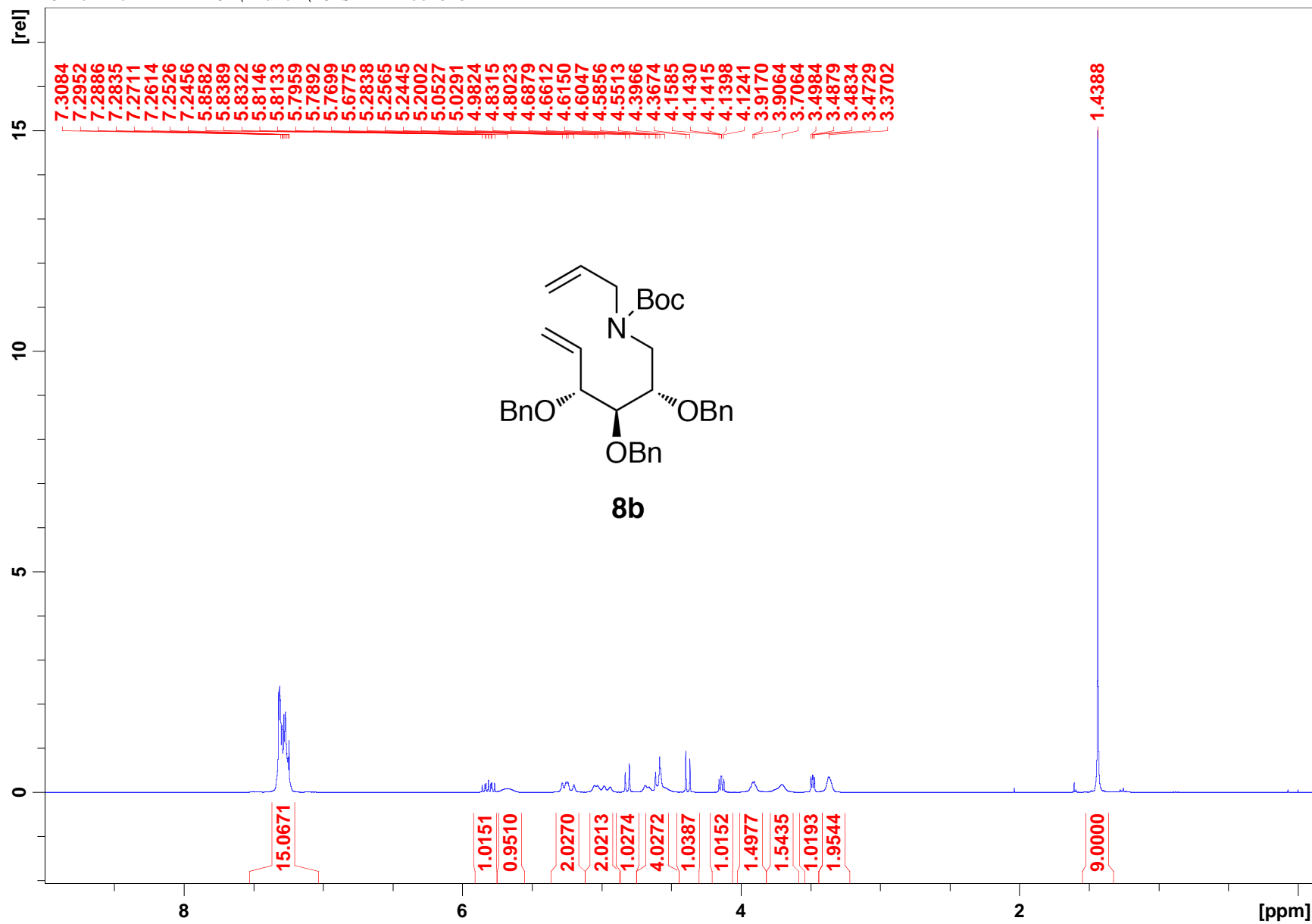
13C-NMR

JL-062207 9 1 C:\Bruker\TOPSPIN leex3451



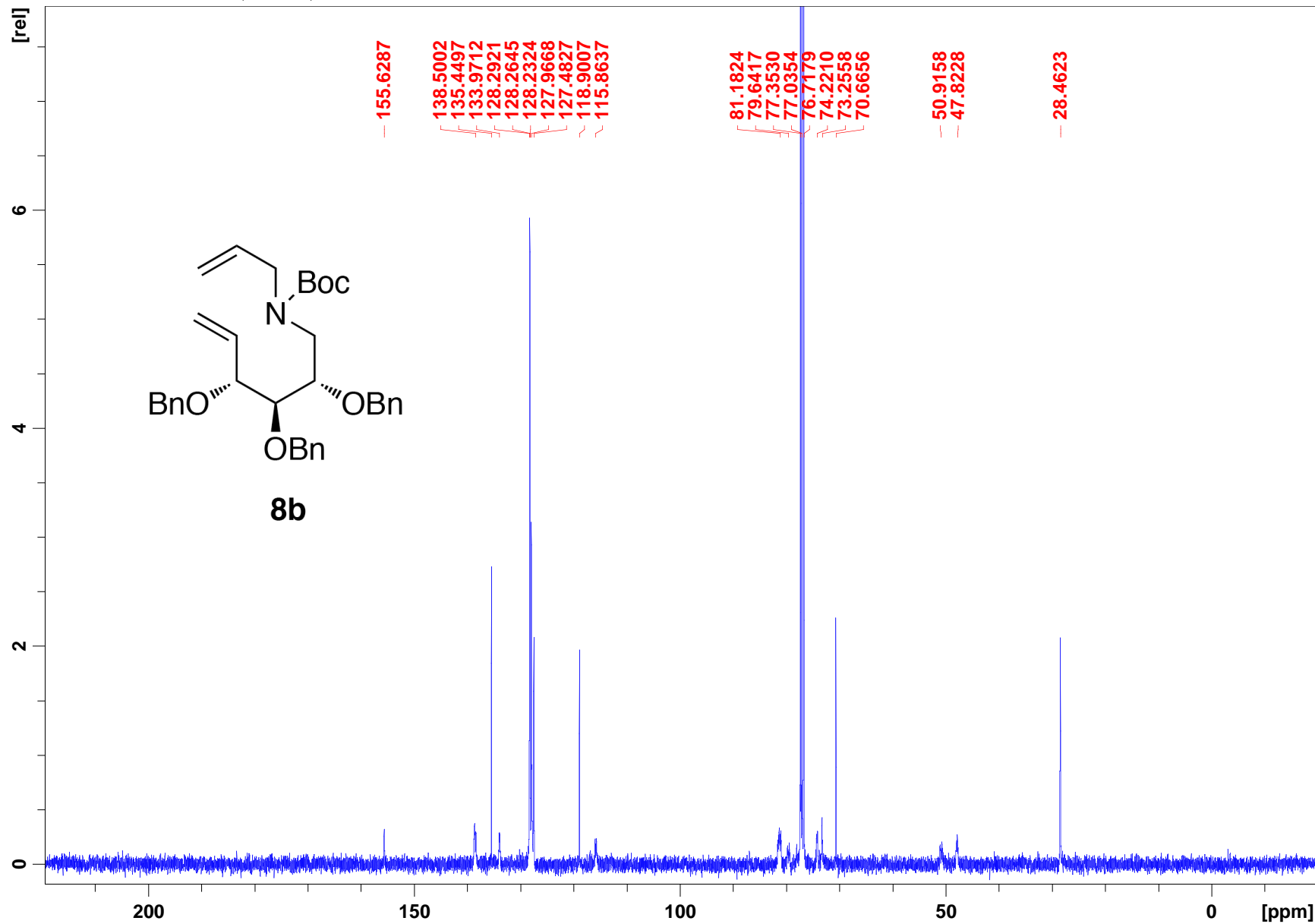
1H-NMR

JL-042707 1 1 C:\Bruker\TOPSPIN leex3451



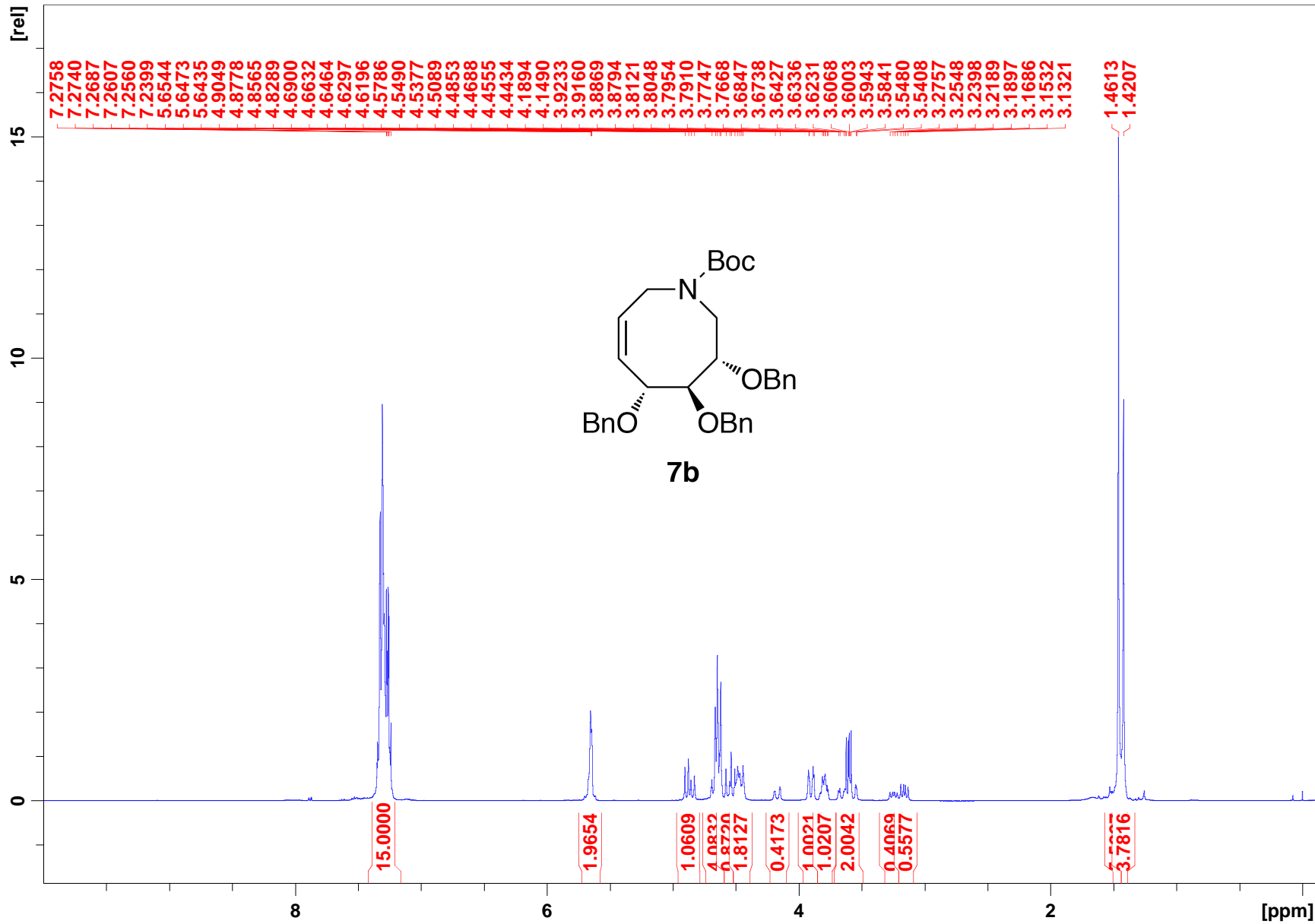
13C-NMR

JL-042707 4 1 C:\Bruker\TOPSPIN leex3451



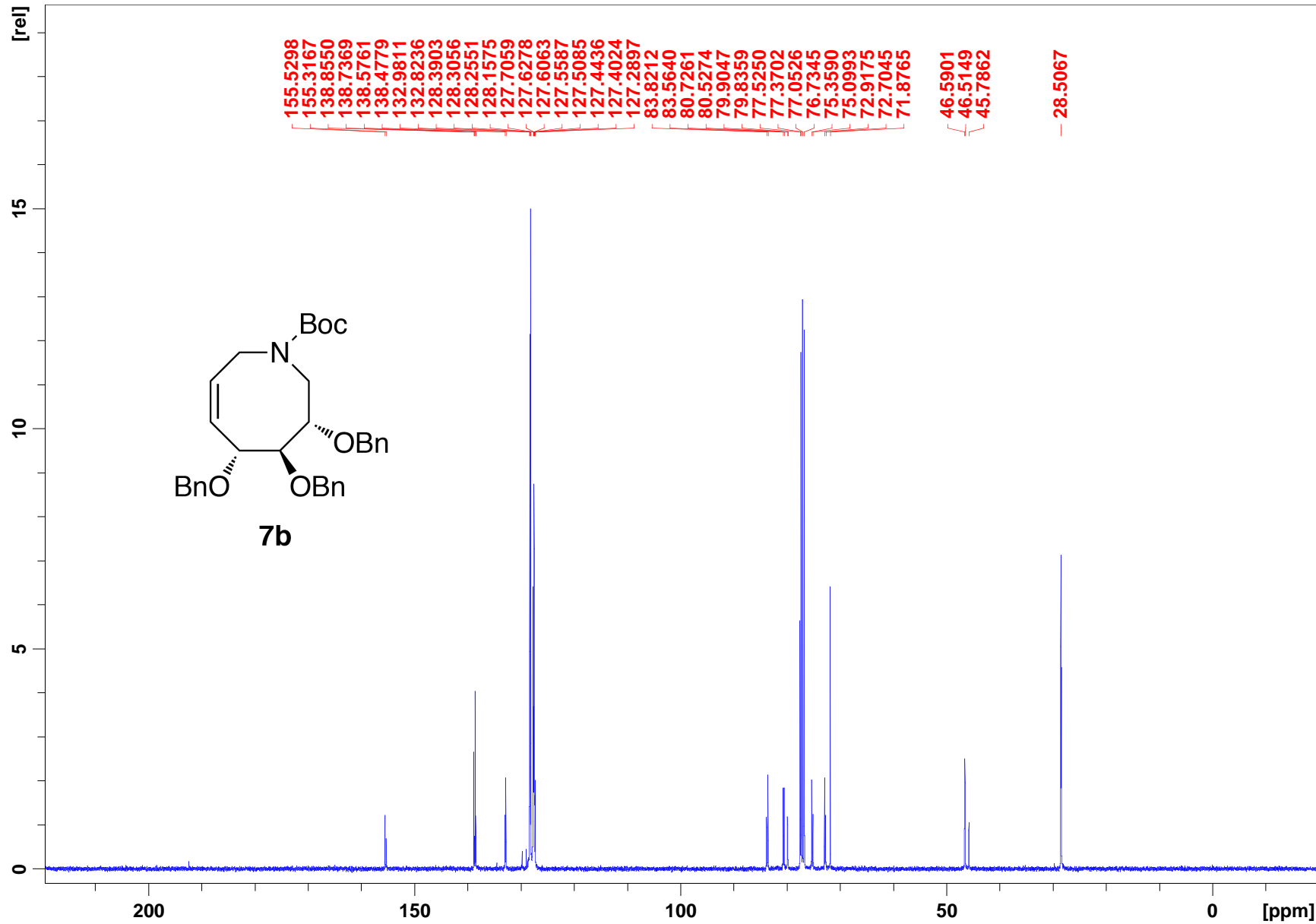
1H-NMR

JL-051807-RE 2 1 C:\Bruker\TOPSPIN leex3451



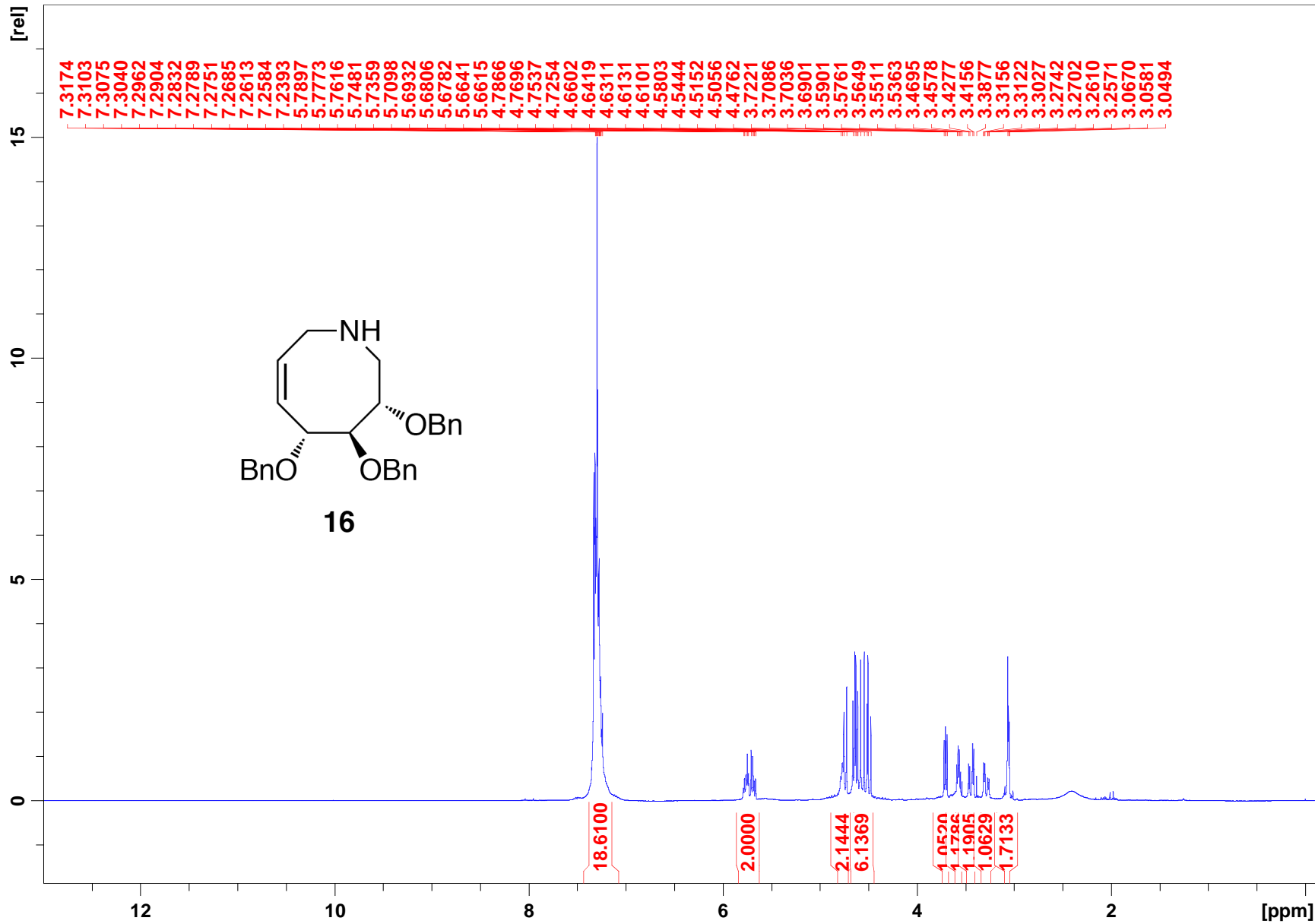
13C-NMR

JL-051807-RE 4 1 C:\Bruker\TOPSPIN leex3451



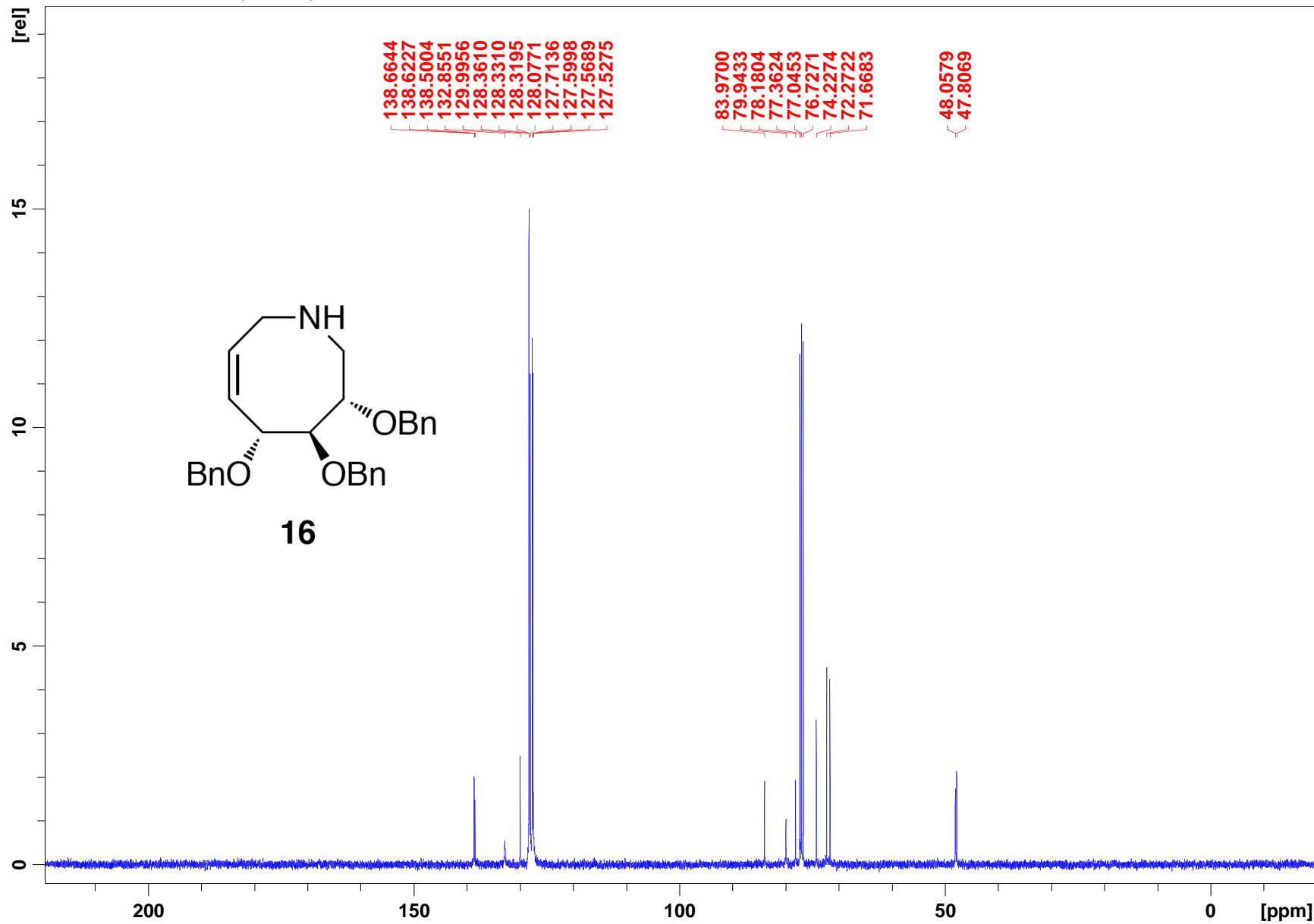
1H-NMR

JL070507 20 1 C:\Bruker\TOPSPIN leex3451



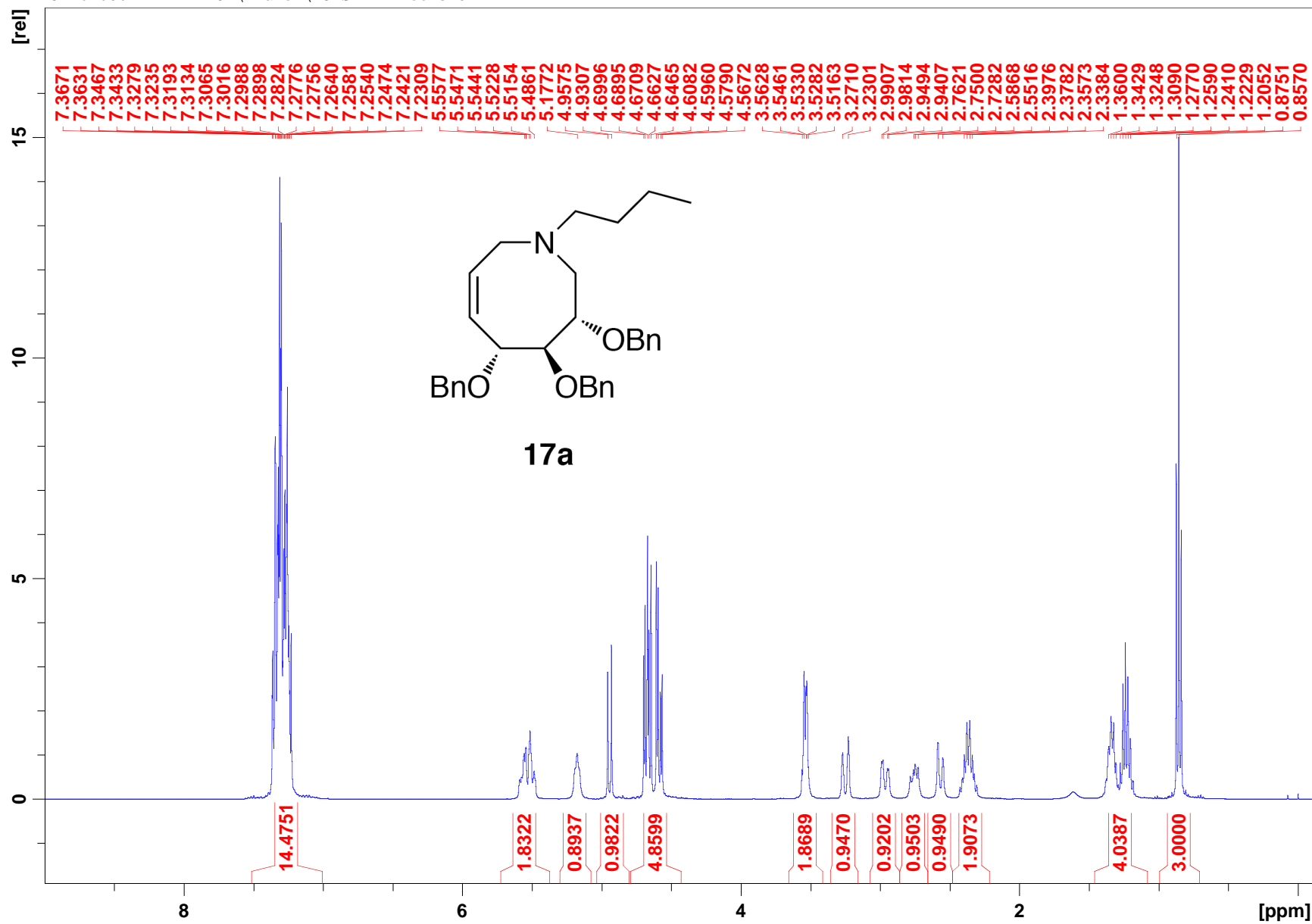
13C-NMR

JL070507 11 1 C:\Bruker\TOPSPIN leex3451



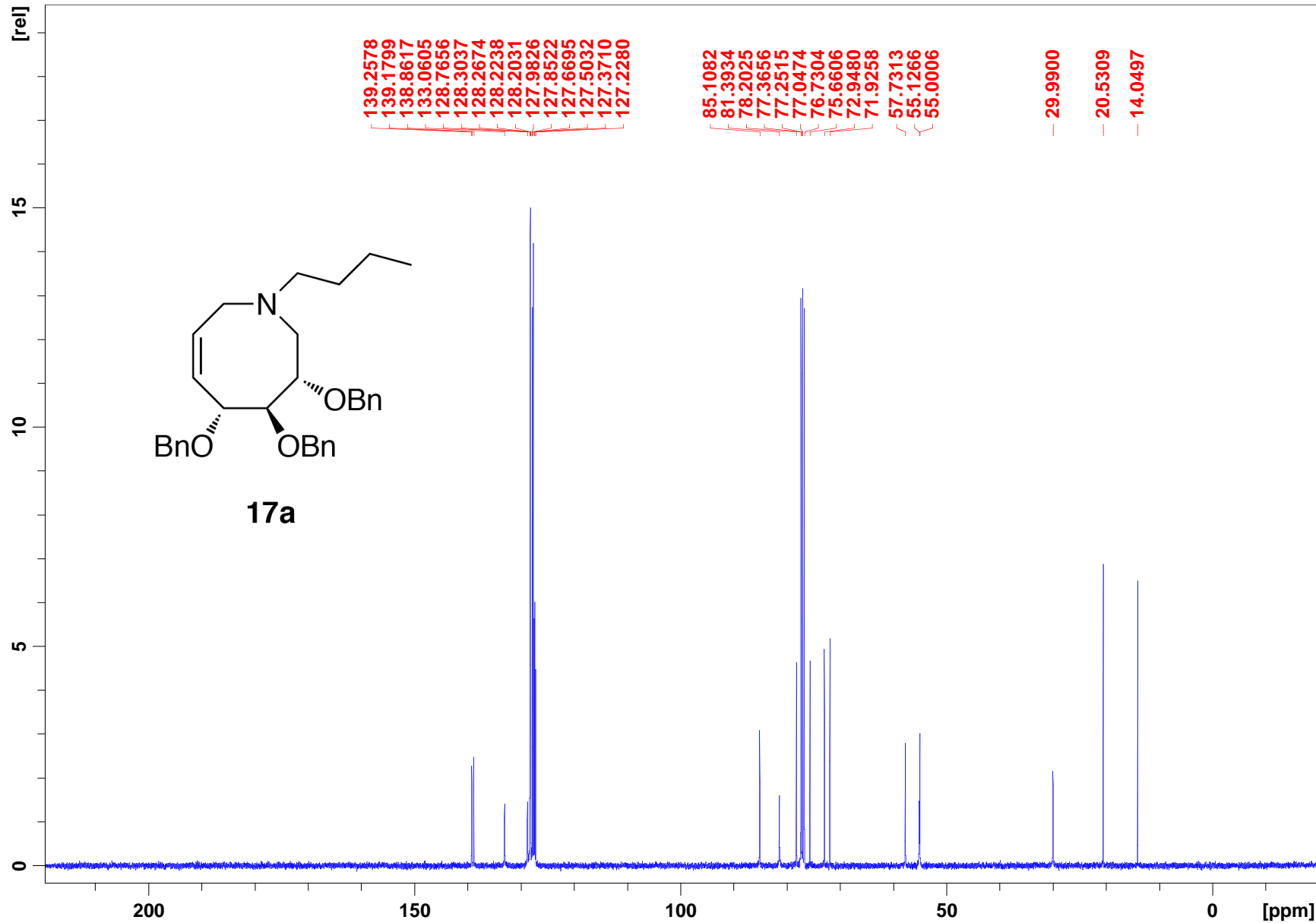
1H-NMR

JL-070807 1 1 C:\Bruker\TOPSPIN leex3451



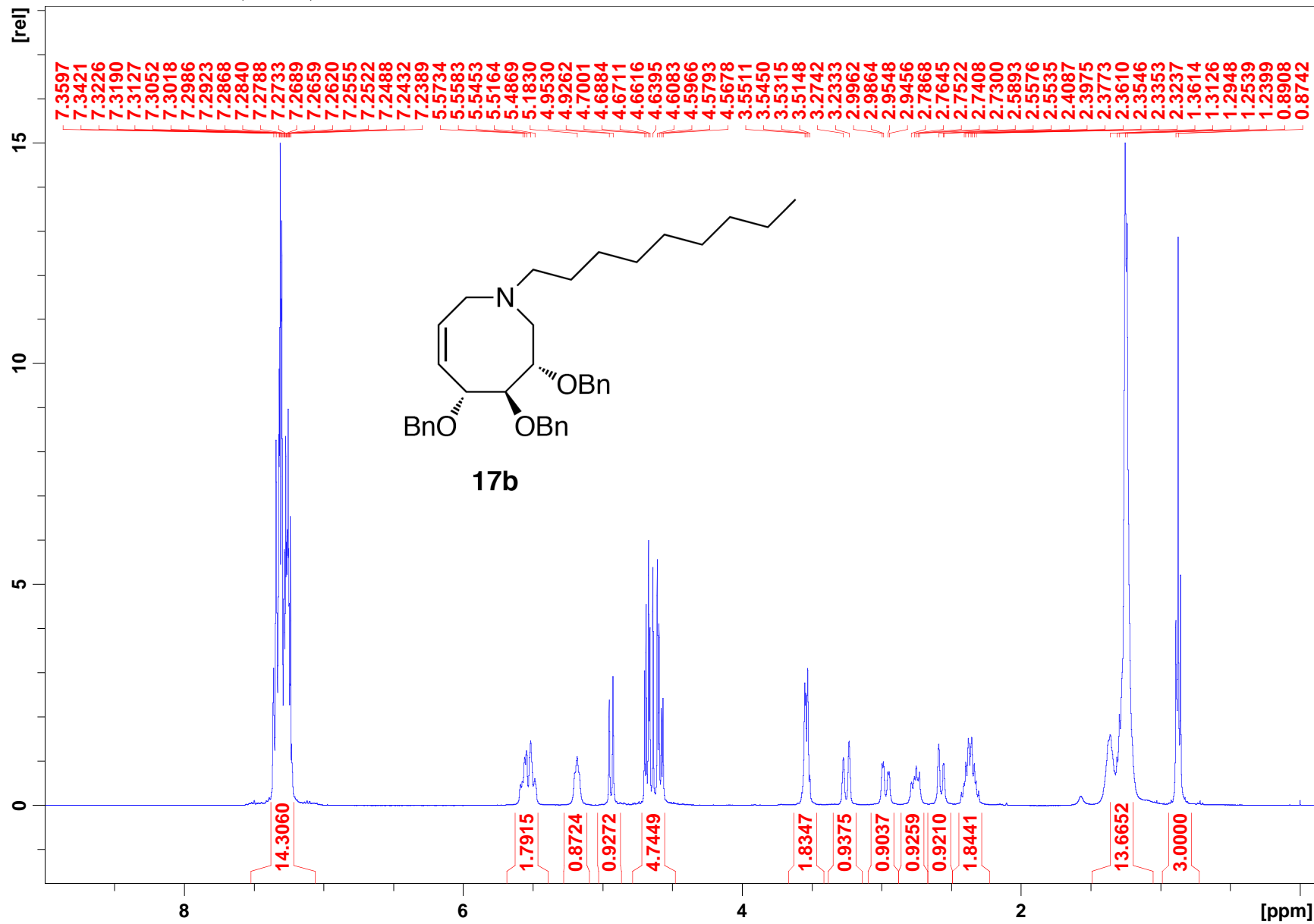
13C-NMR

JL-070807 2 1 C:\Bruker\TOPSPIN leex3451



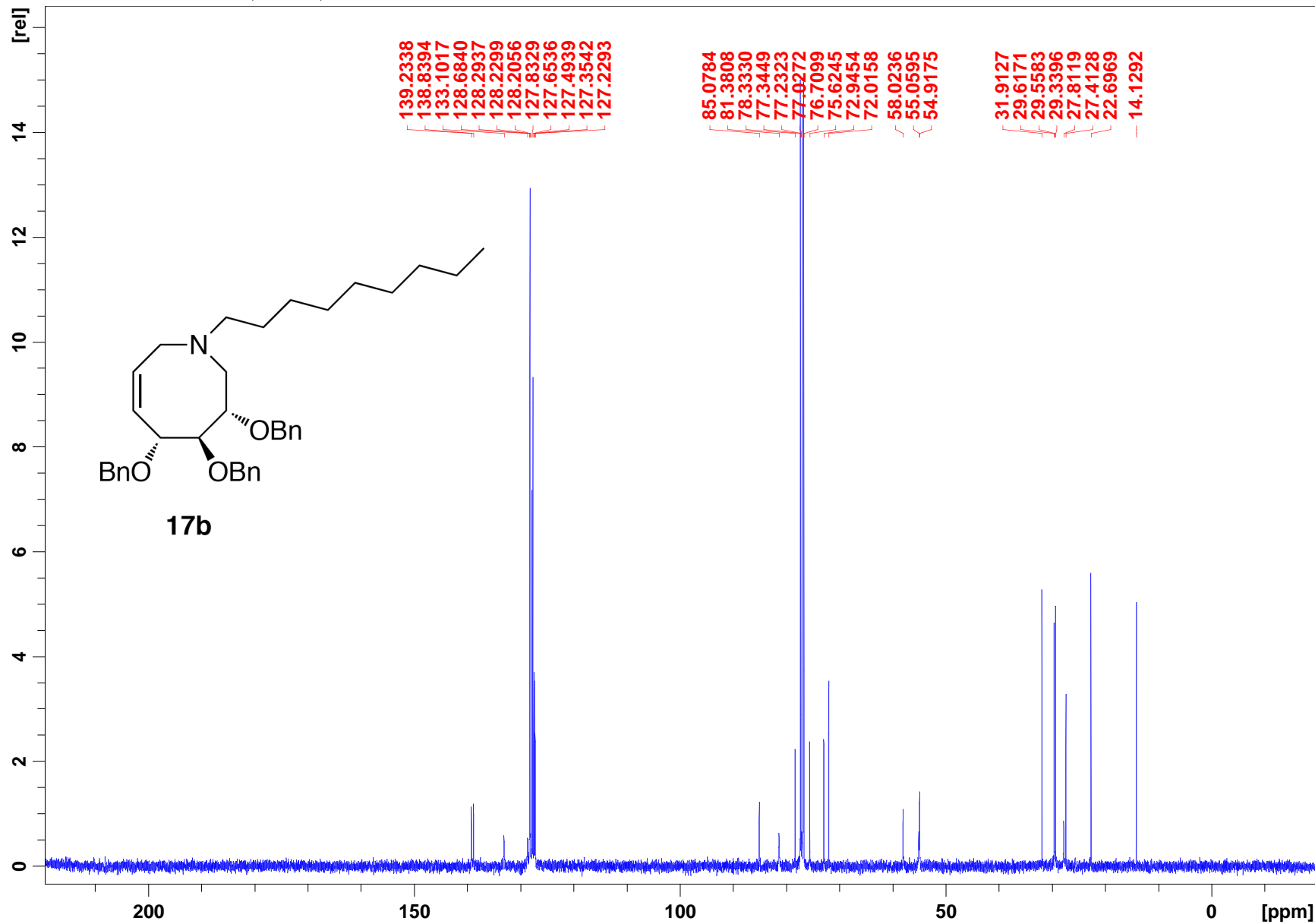
1H-NMR

JL-071307 1 1 C:\Bruker\TOPSPIN leex3451



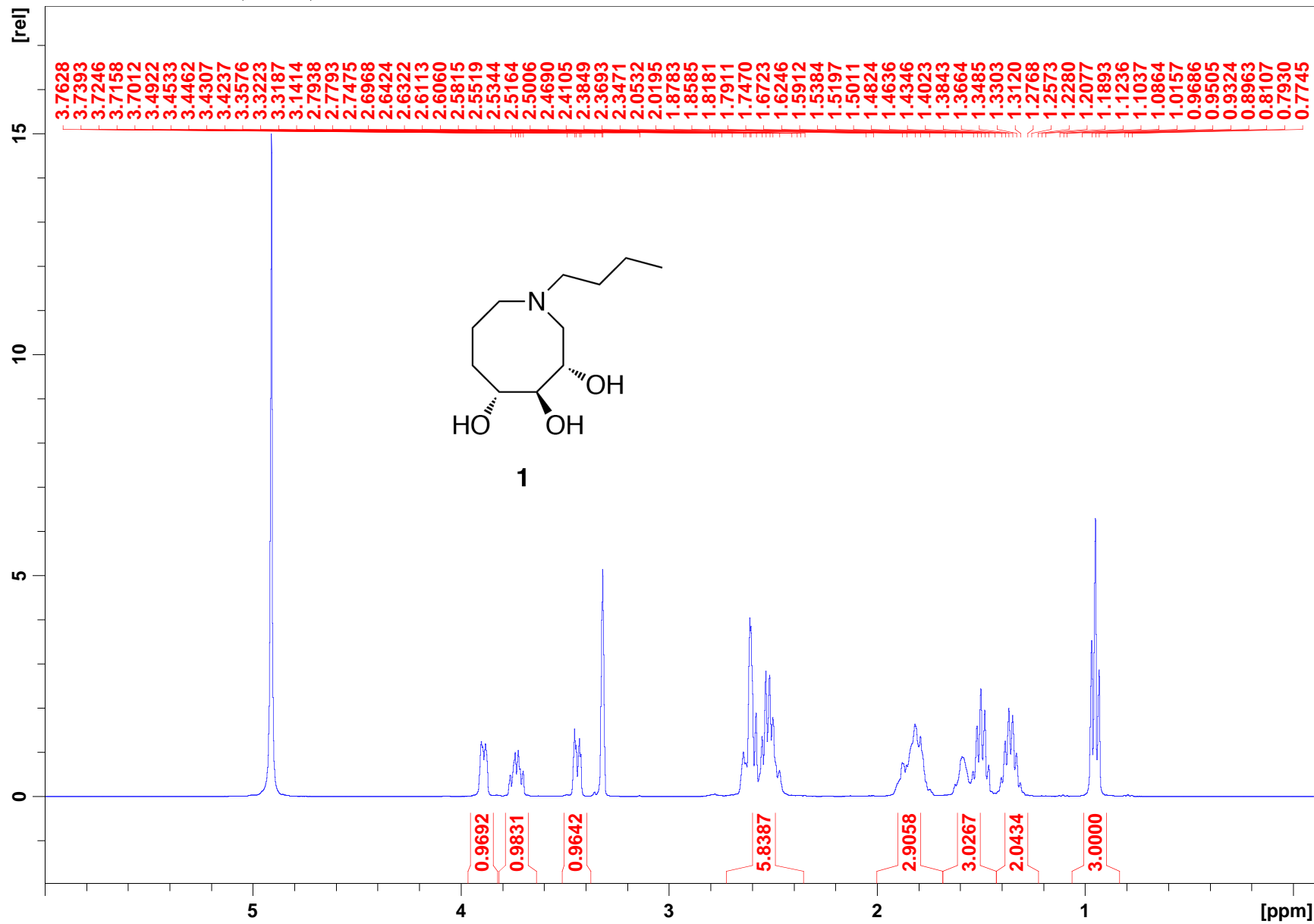
13C-NMR

JL-071307 10 1 C:\Bruker\TOPSPIN leex3451



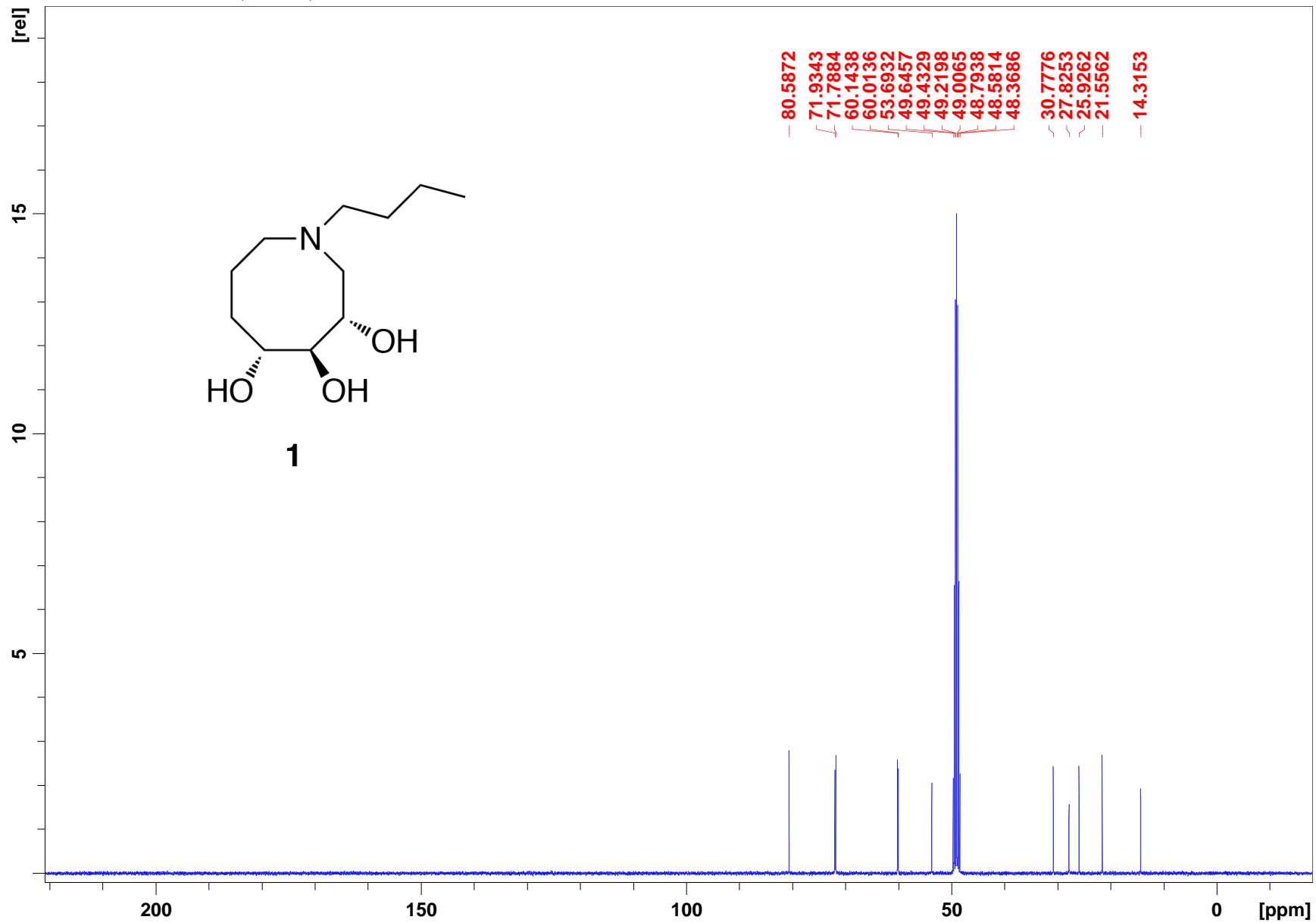
1H-NMR

JL-030608 1 1 C:\Bruker\TOPSPIN leex3451



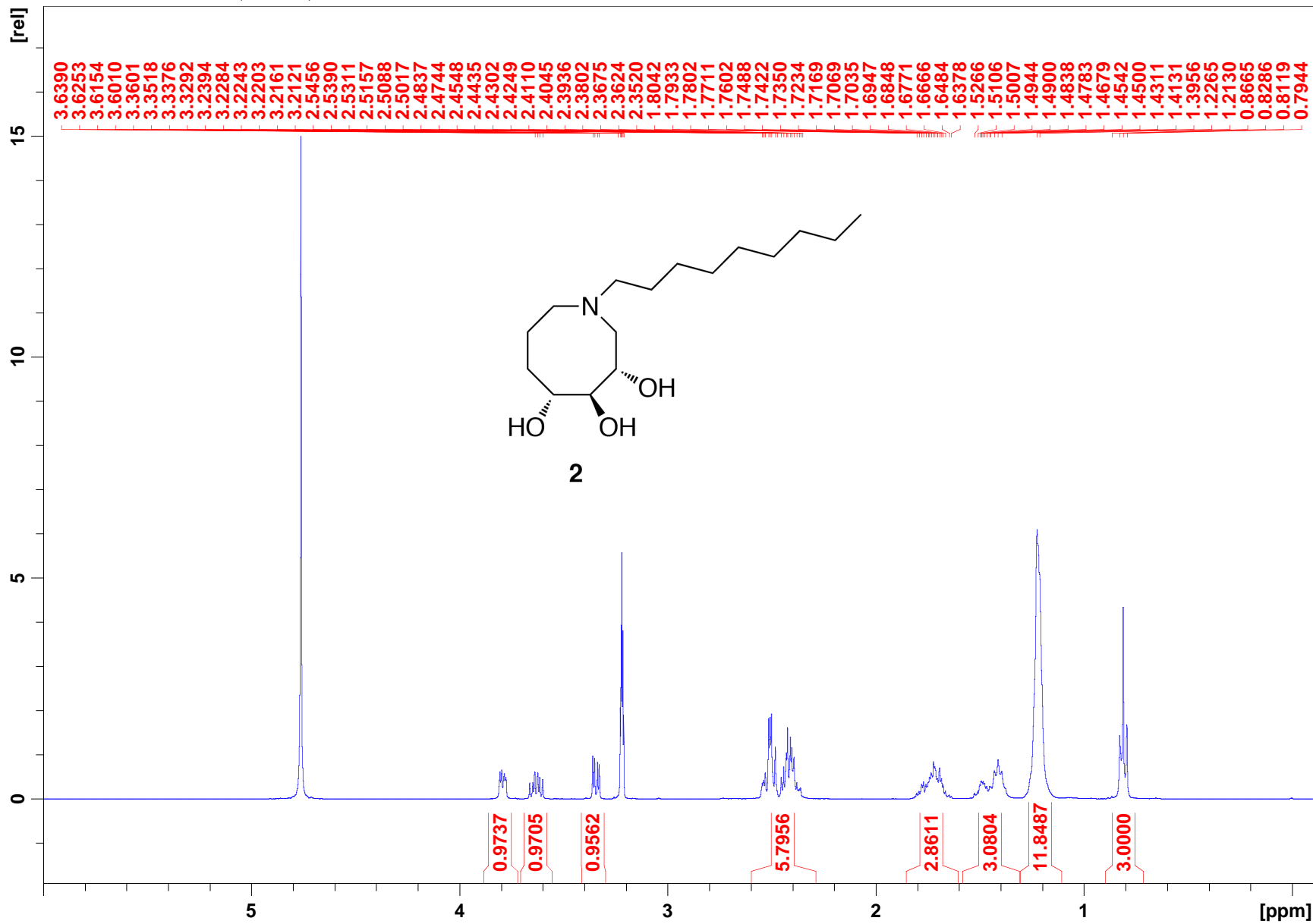
13C-NMR

JL-030608 3 1 C:\Bruker\TOPSPIN leex3451



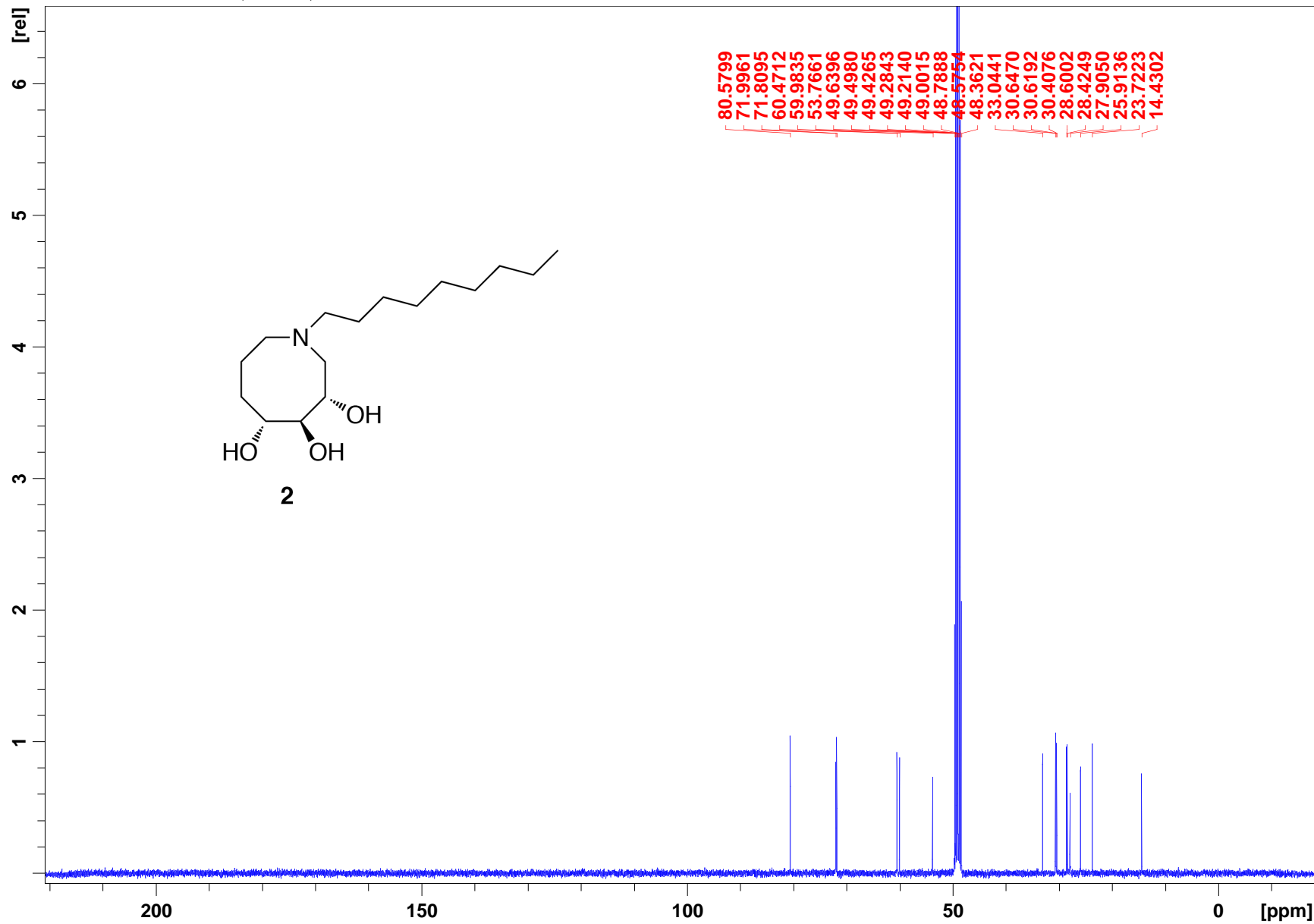
1H-NMR

JL-030708 1 1 C:\Bruker\TOPSPIN leex3451



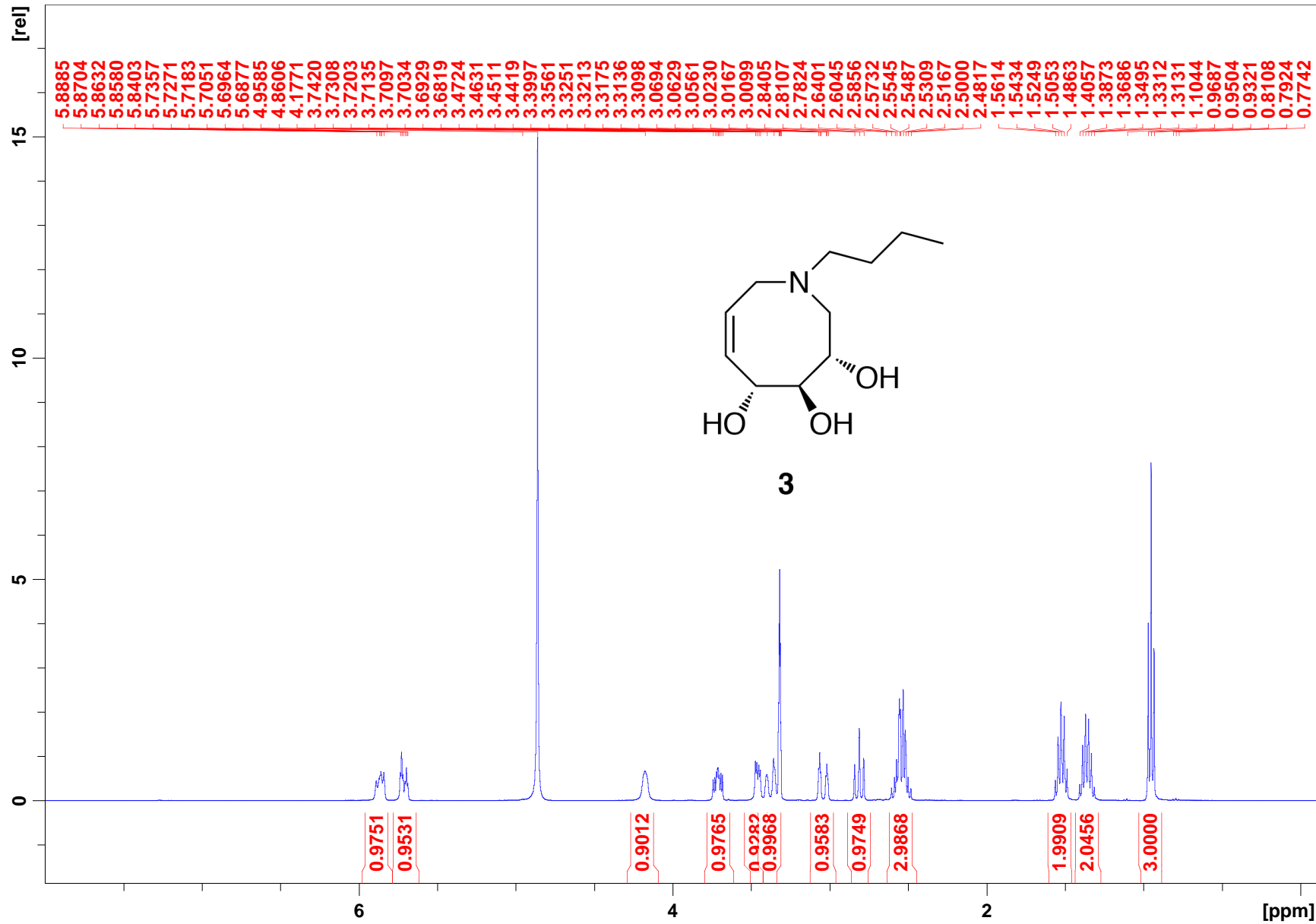
13C-NMR

JL-030708 3 1 C:\Bruker\TOPSPIN leex3451



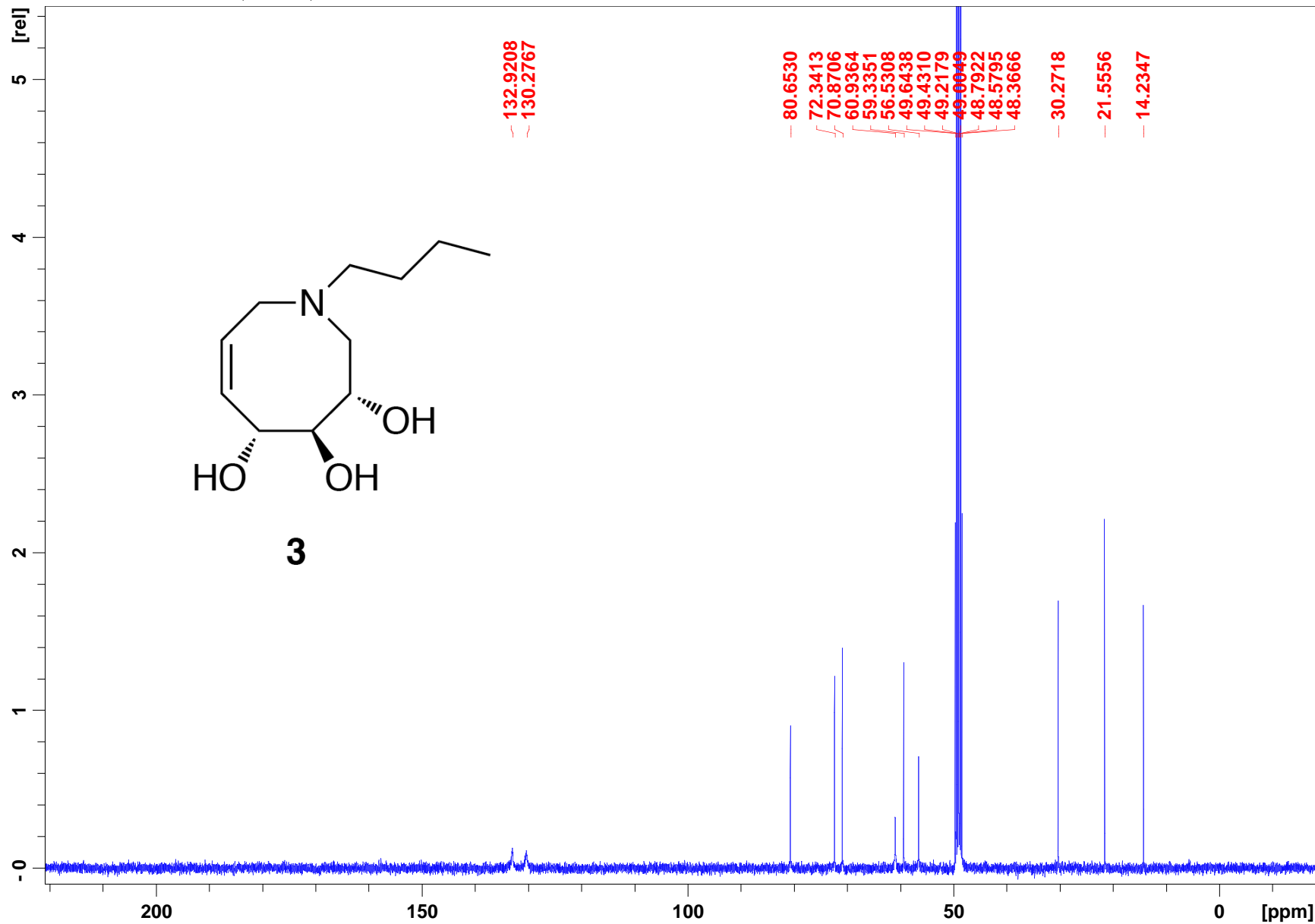
1H-NMR

JL-032008 1 1 C:\Bruker\TOPSPIN leex3451



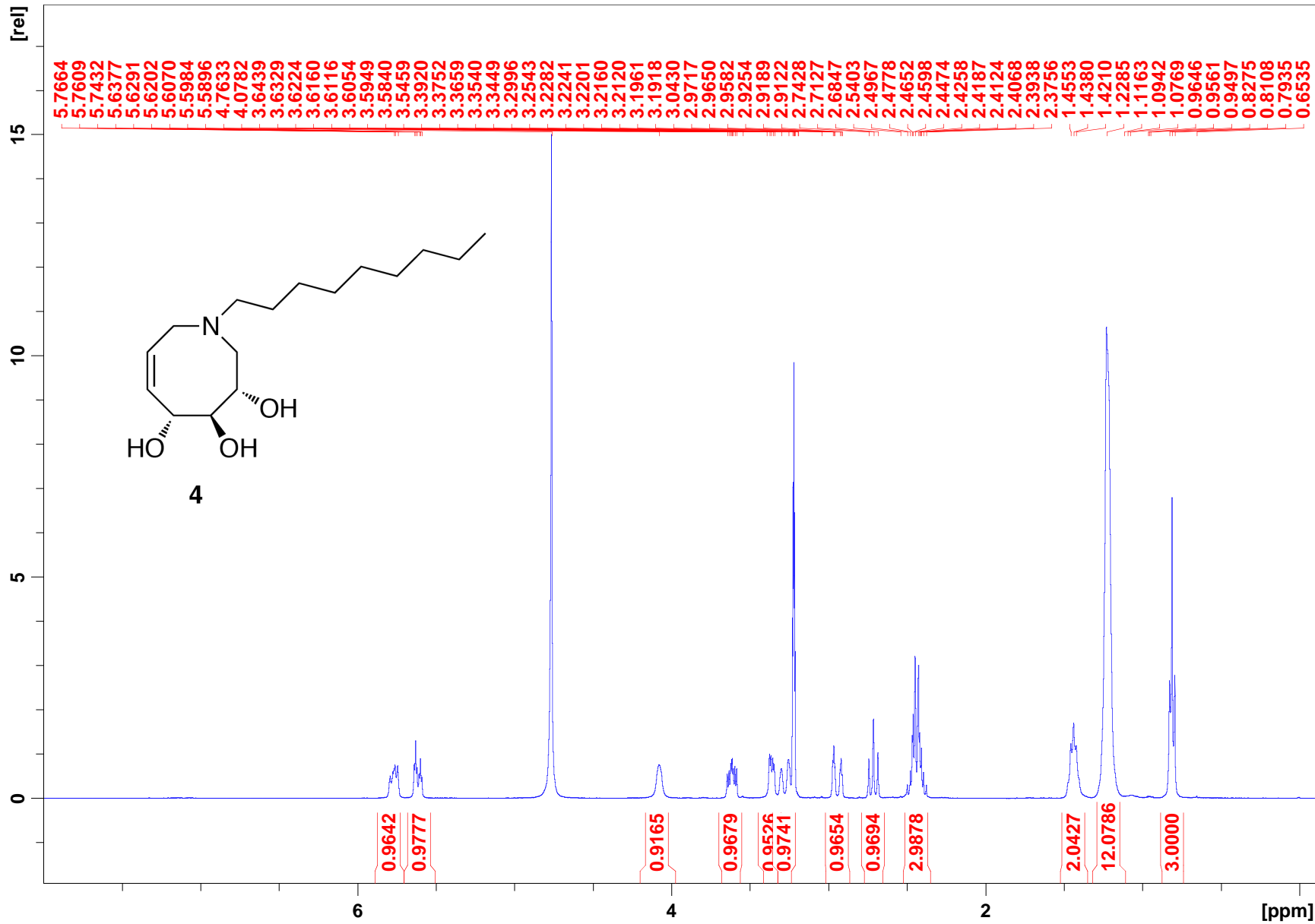
13C-NMR

JL-032008 3 1 C:\Bruker\TOPSPIN leex3451



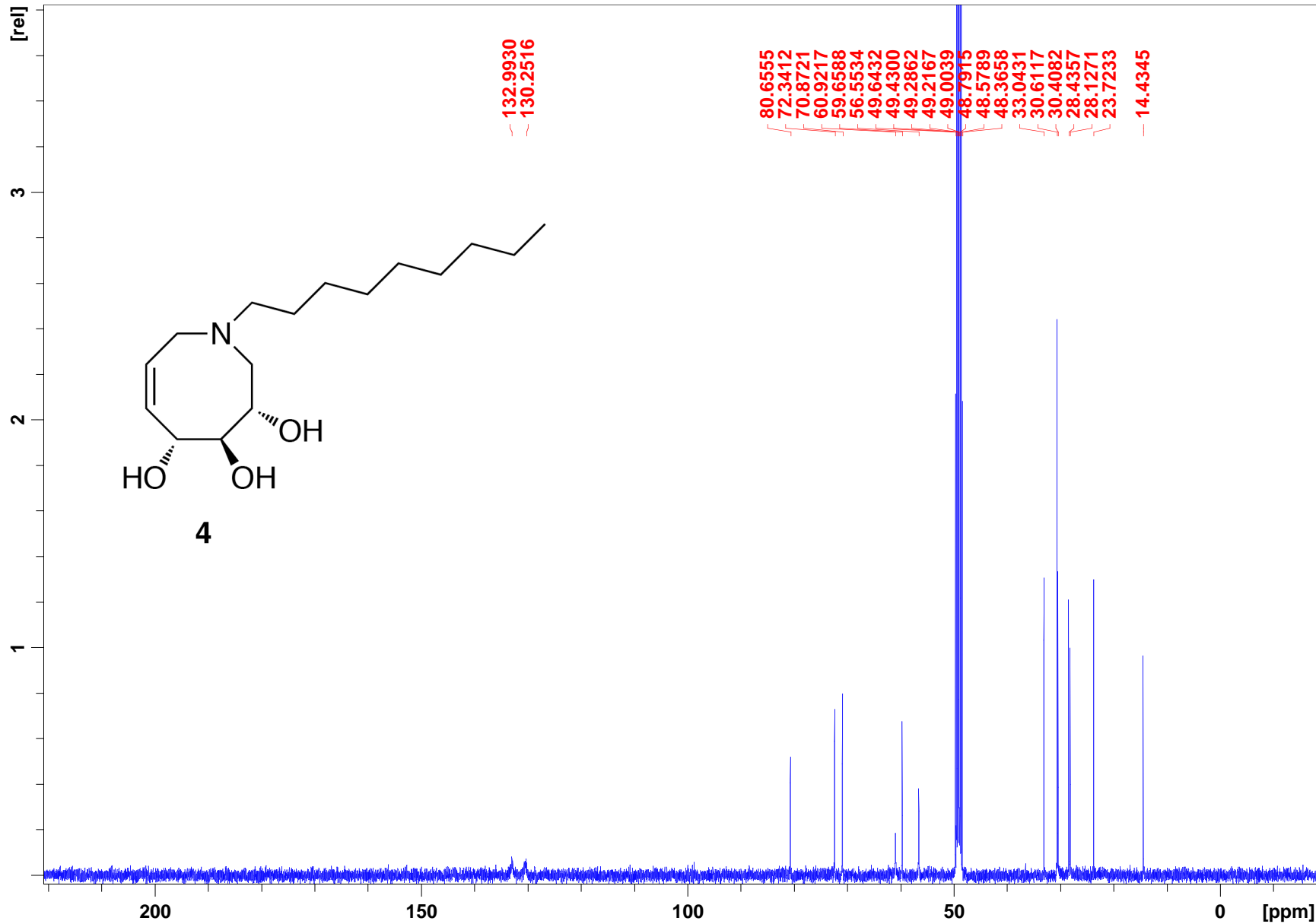
1H-NMR

JL-031908-1 1 1 C:\Bruker\TOPSPIN leex3451



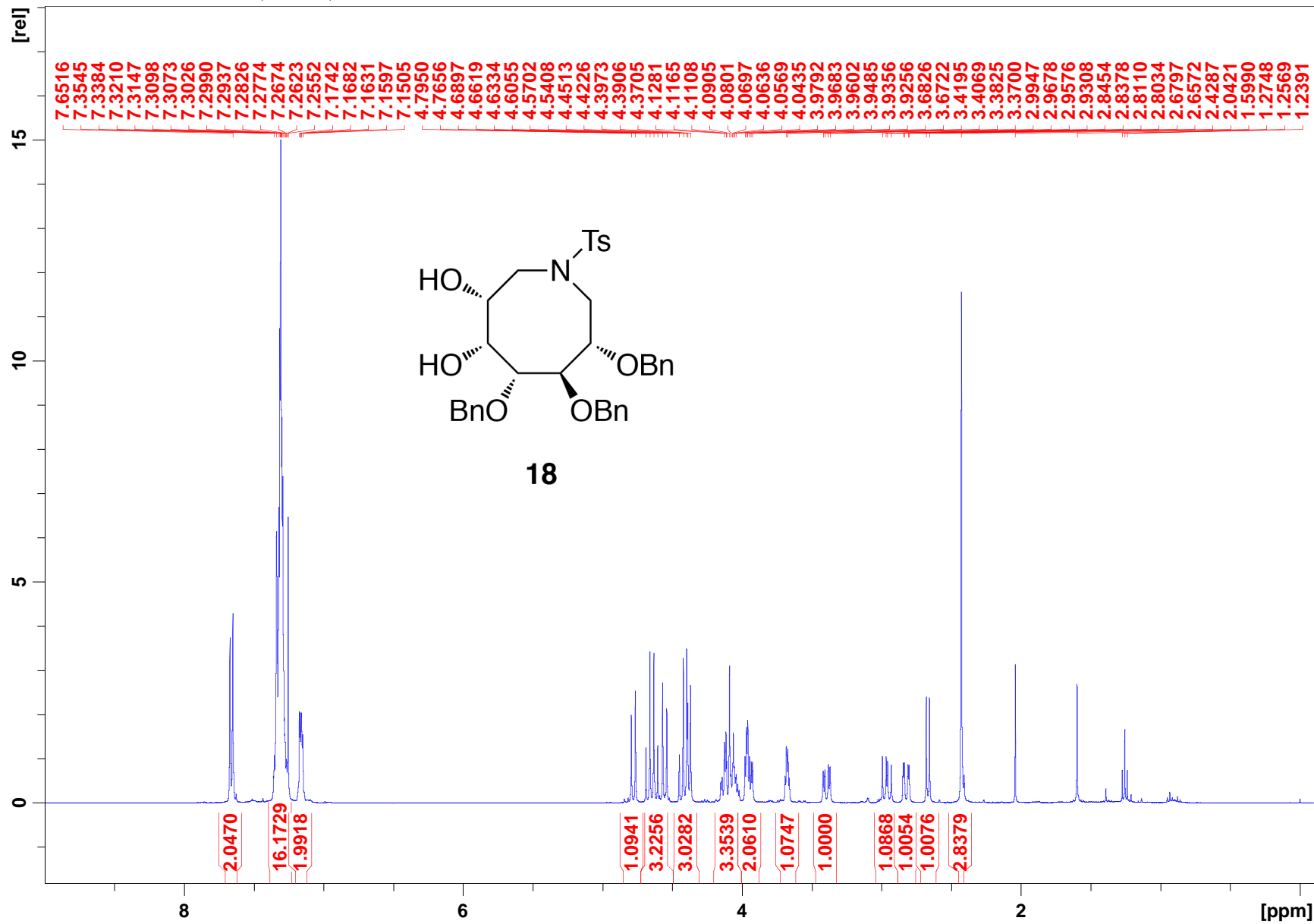
13C-NMR

JL-031908 3 1 C:\Bruker\TOPSPIN leex3451



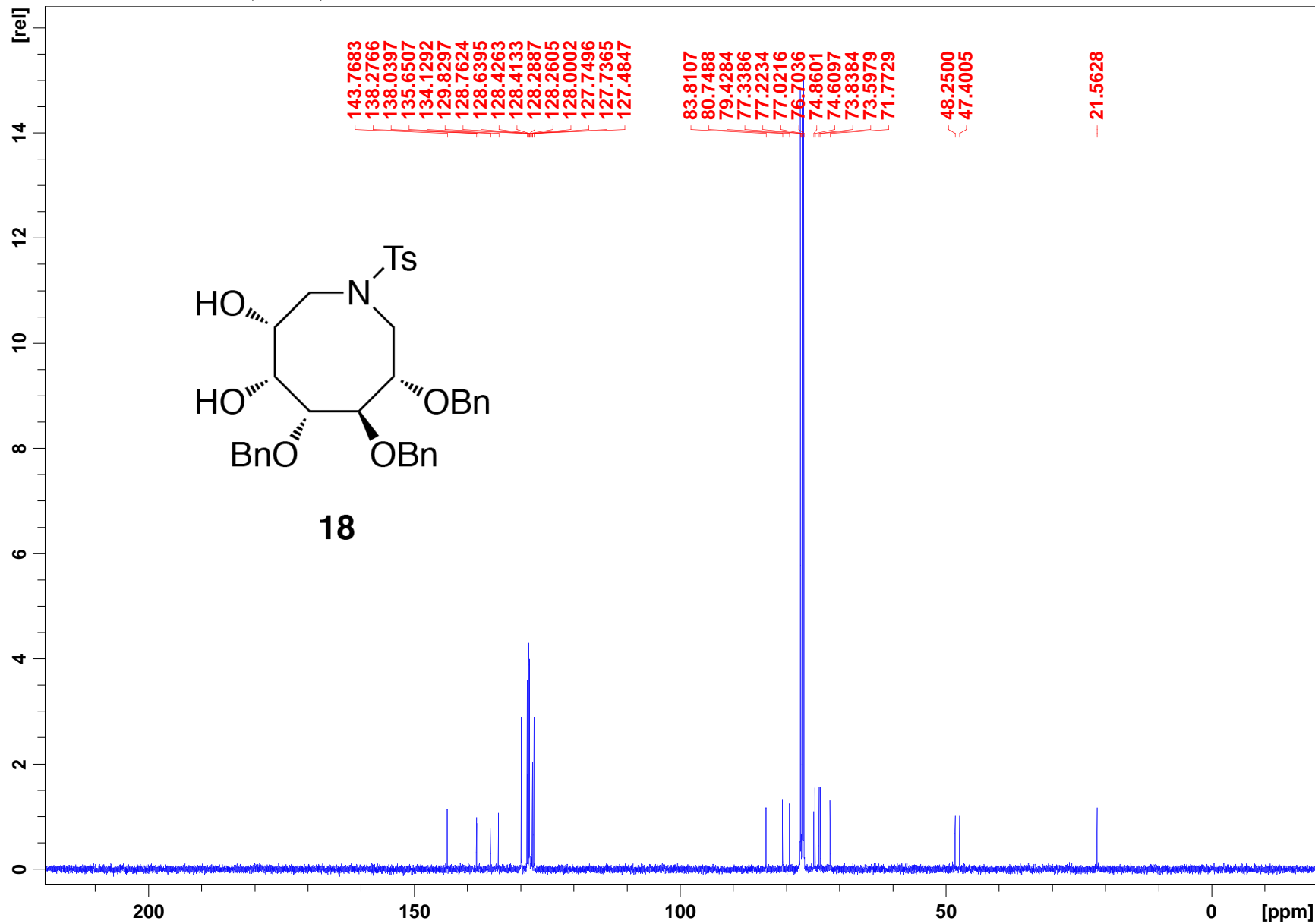
1H-NMR

JL-110107-B 1 1 C:\Bruker\TOPSPIN leex3451



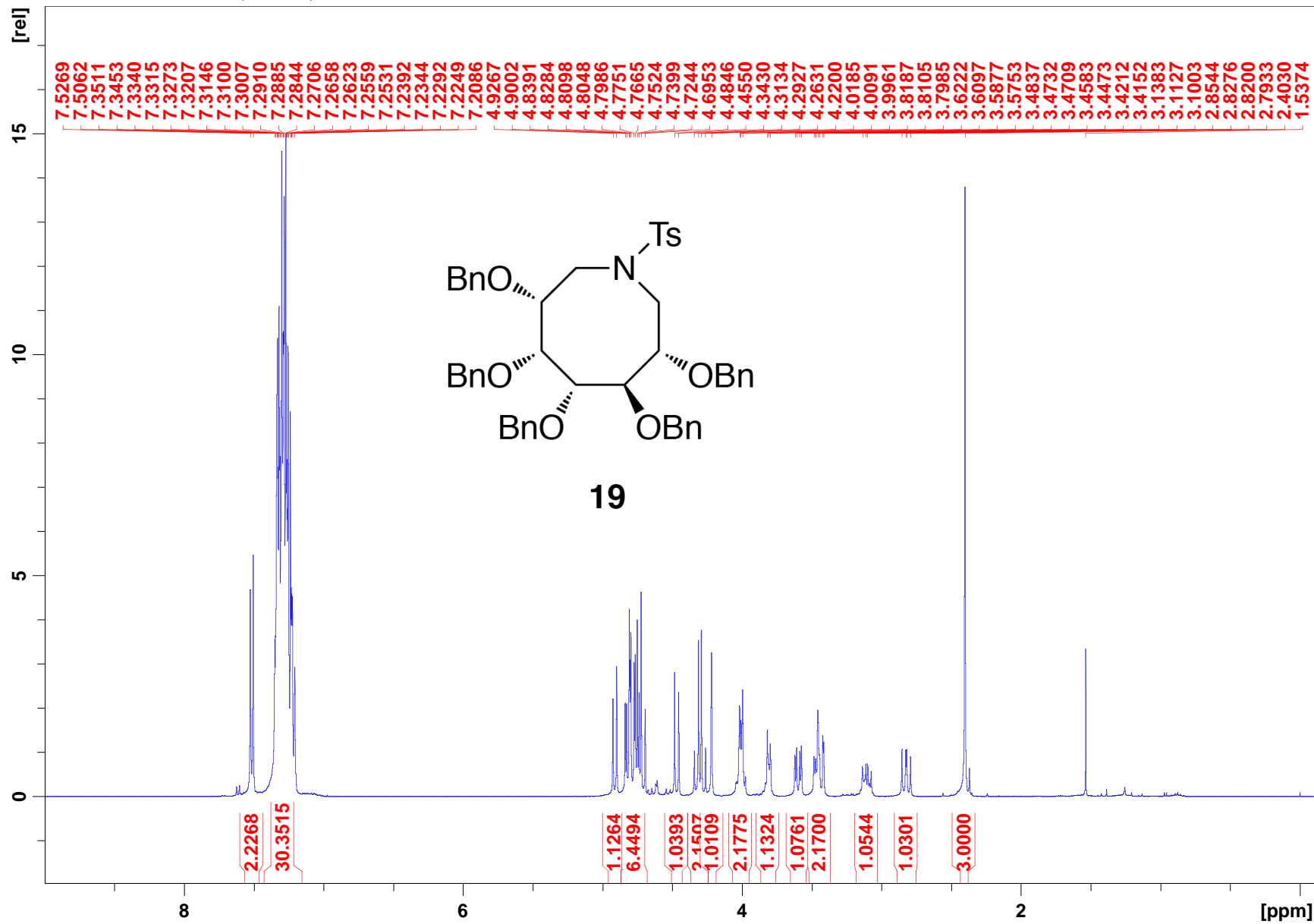
13C-NMR

JL-011208 12 1 C:\Bruker\TOPSPIN leex3451



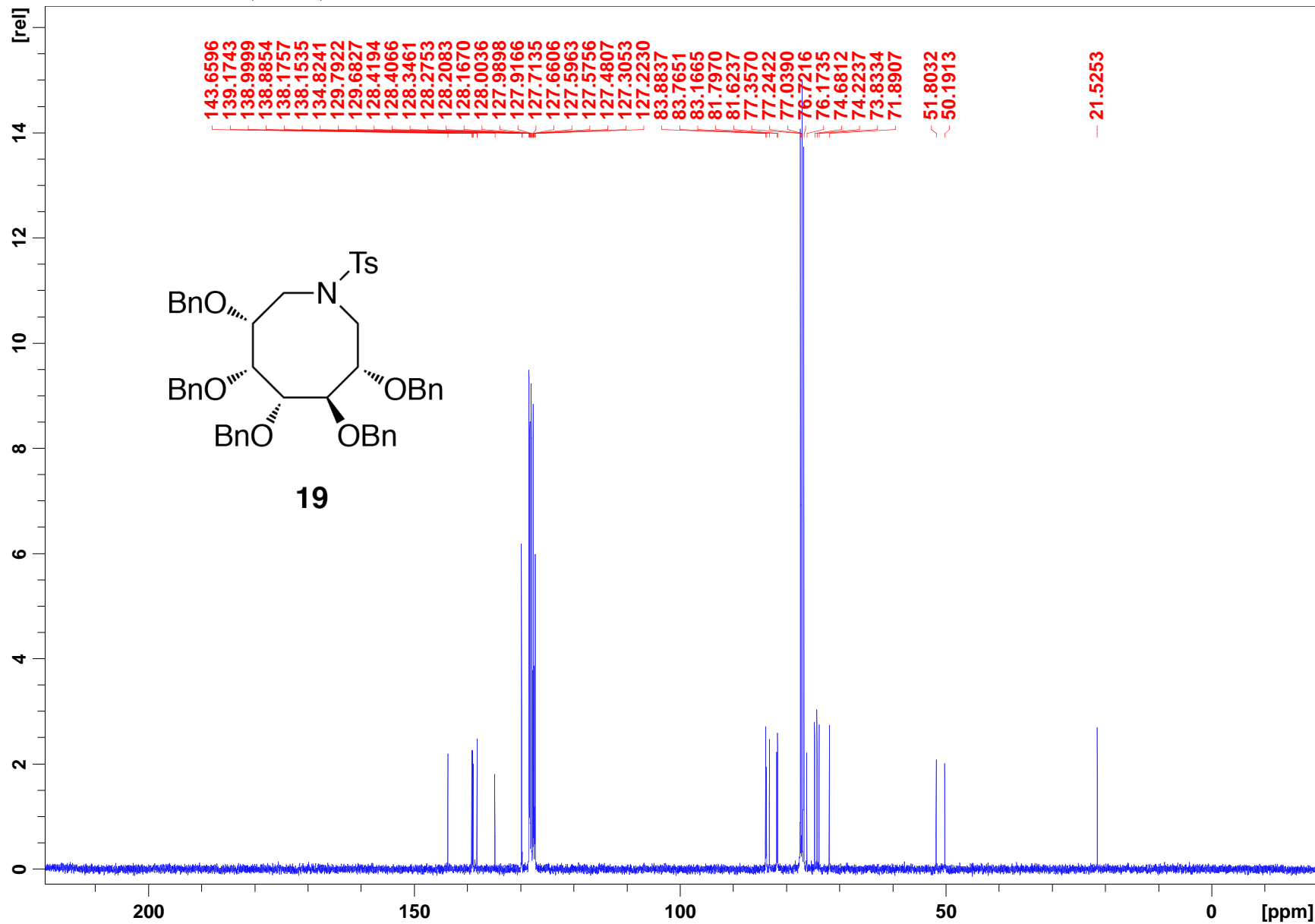
1H-NMR

JL-110807 1 1 C:\Bruker\TOPSPIN leex3451



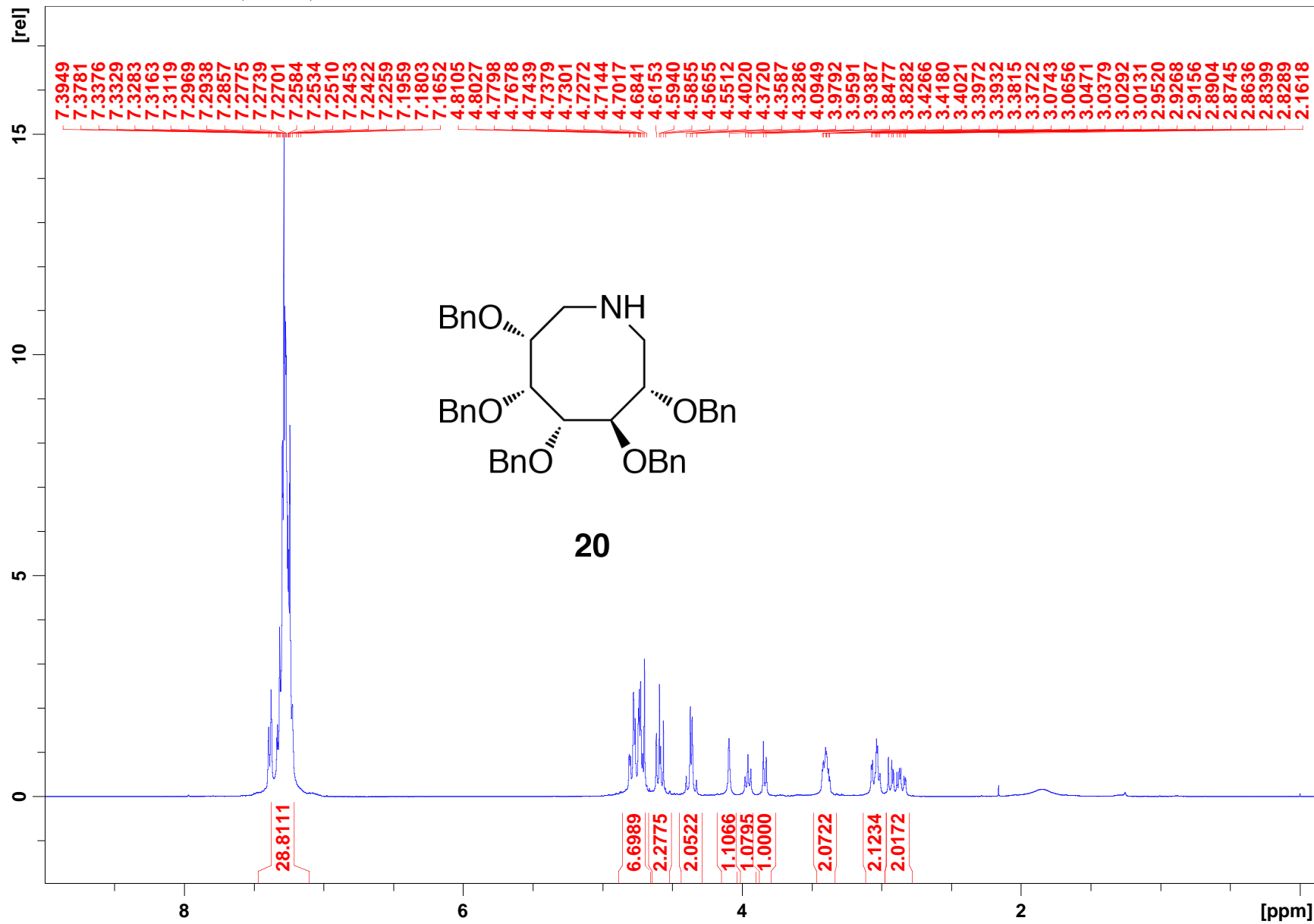
13C-NMR

LJC-110807 4 1 C:\Bruker\TOPSPIN leex3451



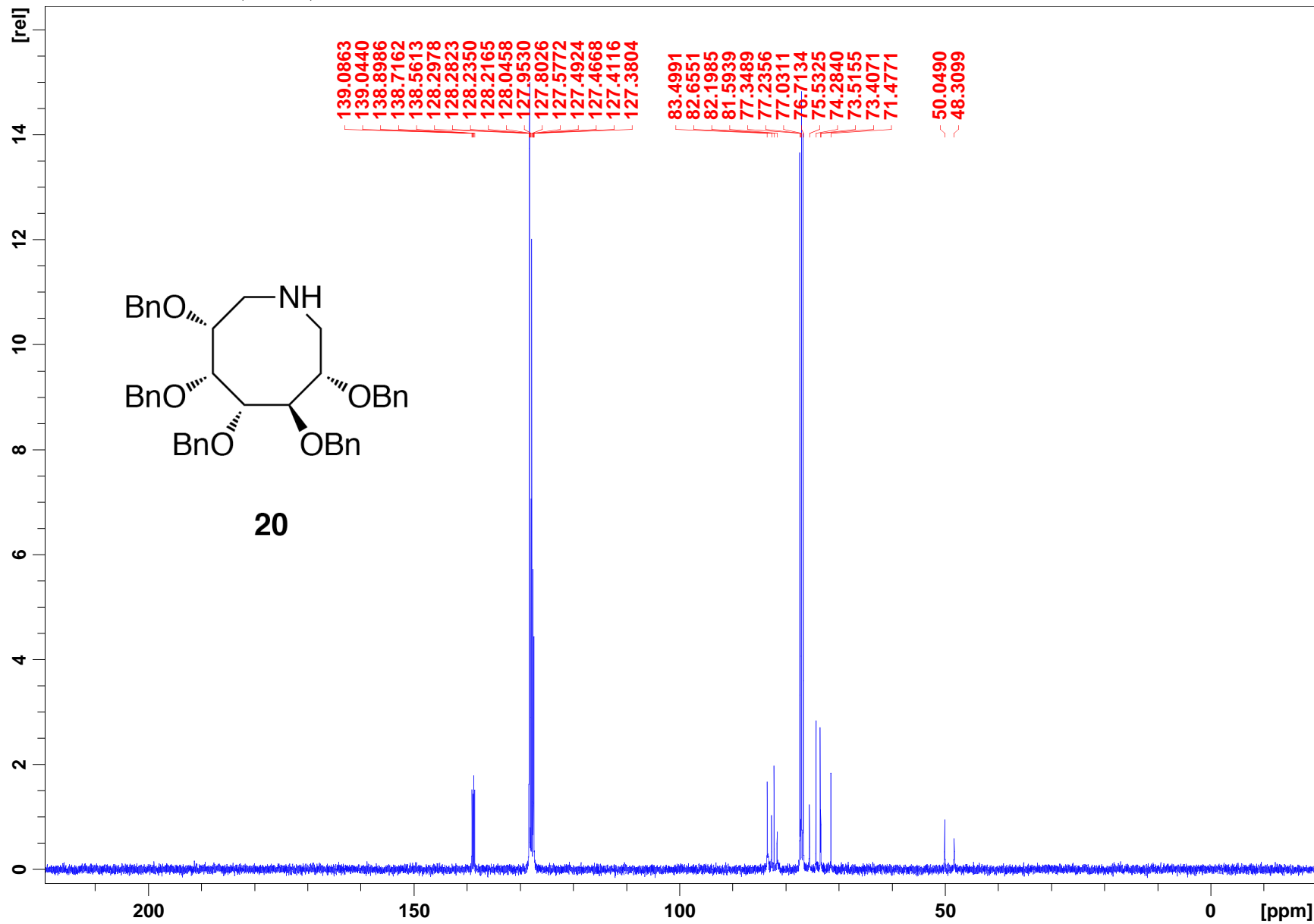
1H-NMR

JL-111907 1 1 C:\Bruker\TOPSPIN leex3451



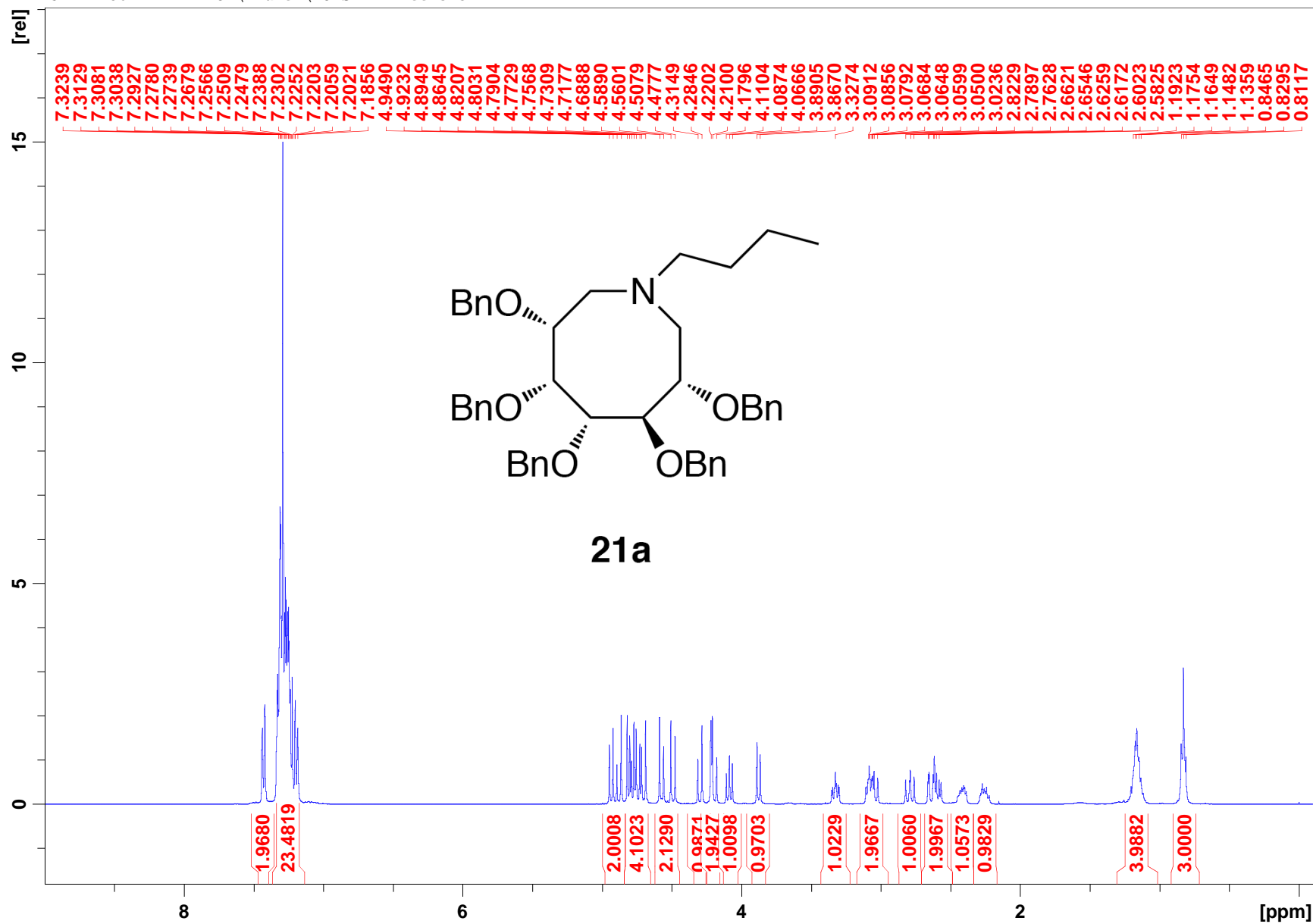
13C-NMR

JL-111907 3 1 C:\Bruker\TOPSPIN leex3451



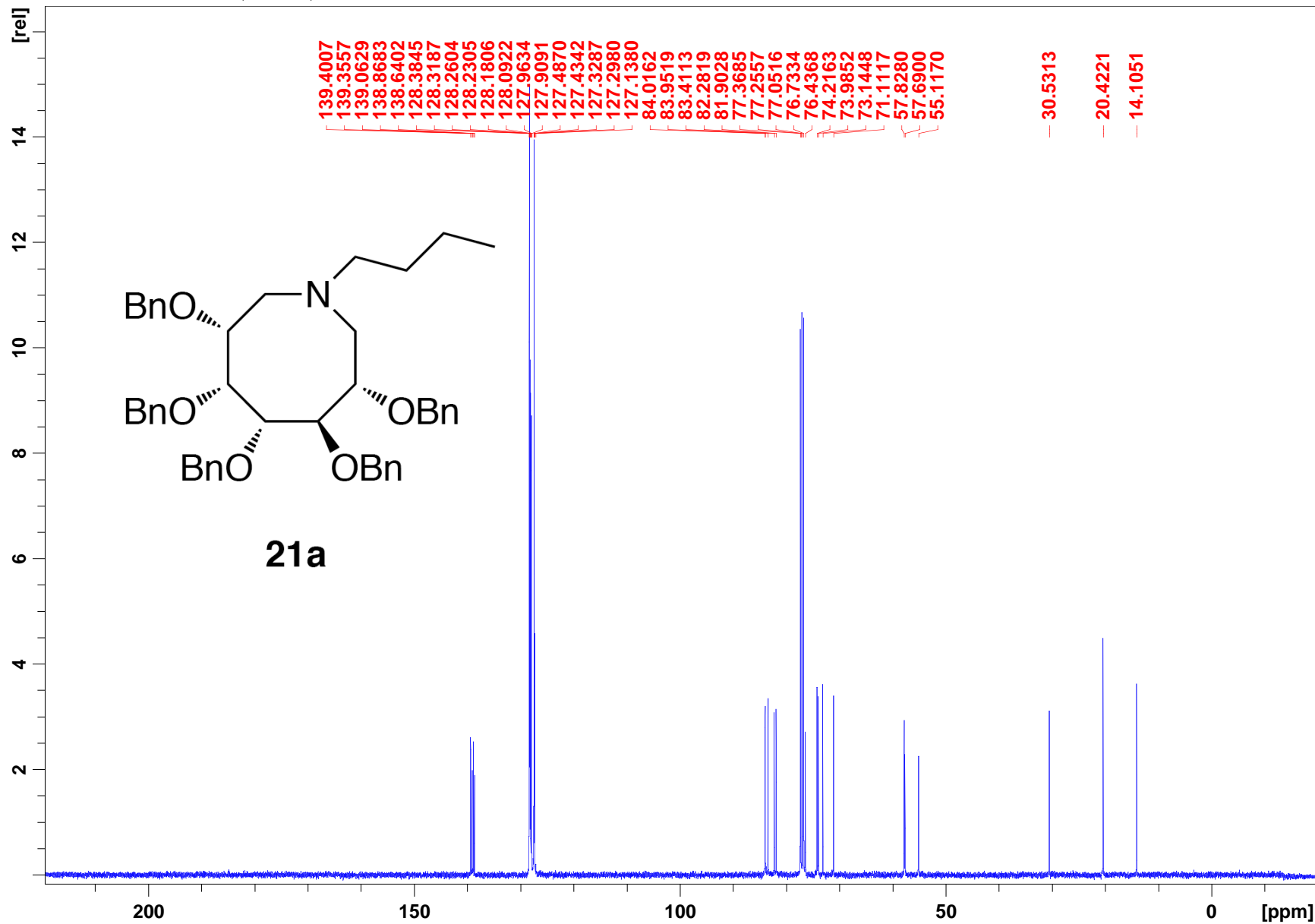
1H-NMR

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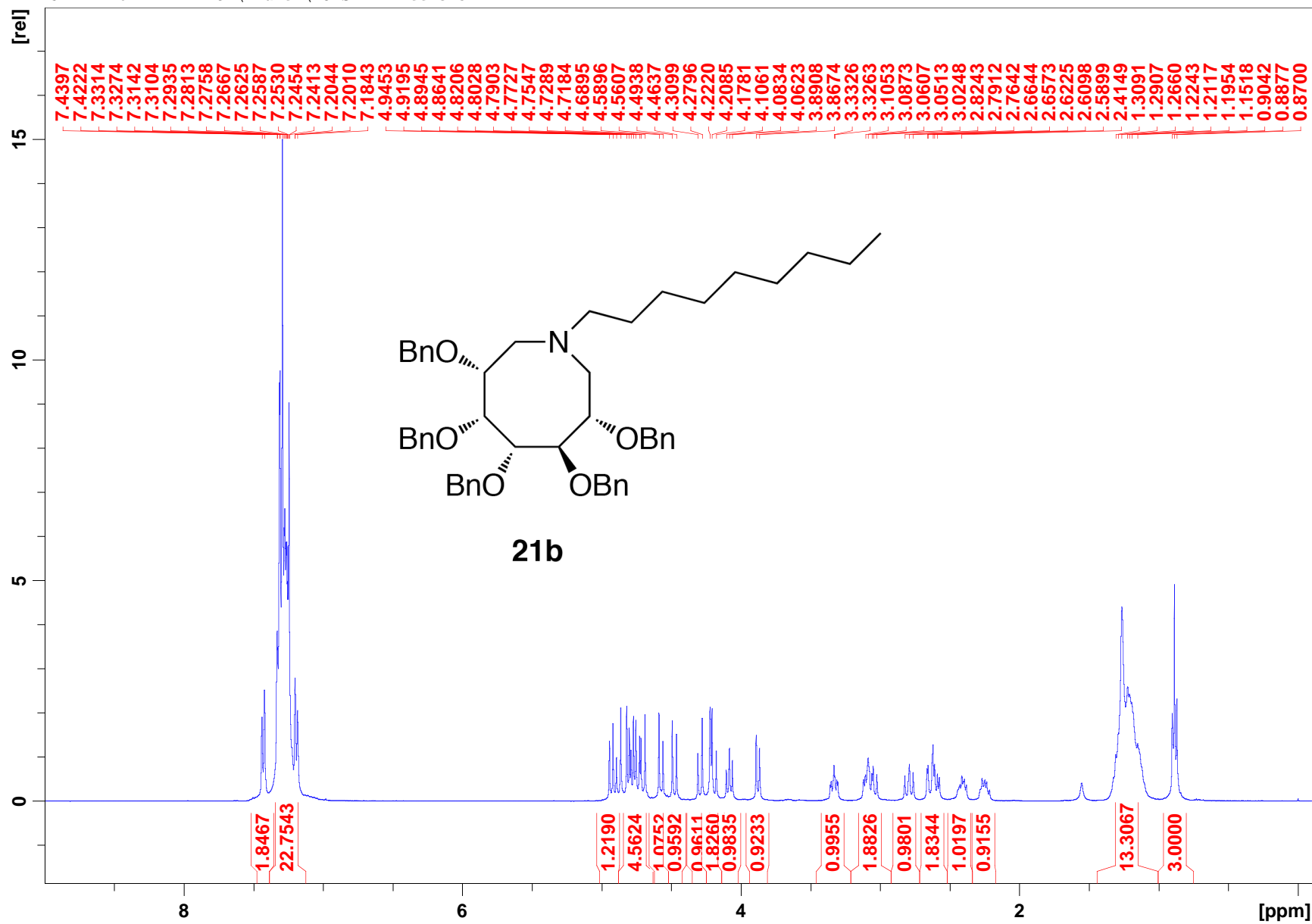
13C-NMR

JL-121807 3 1 C:\Bruker\TOPSPIN leex3451



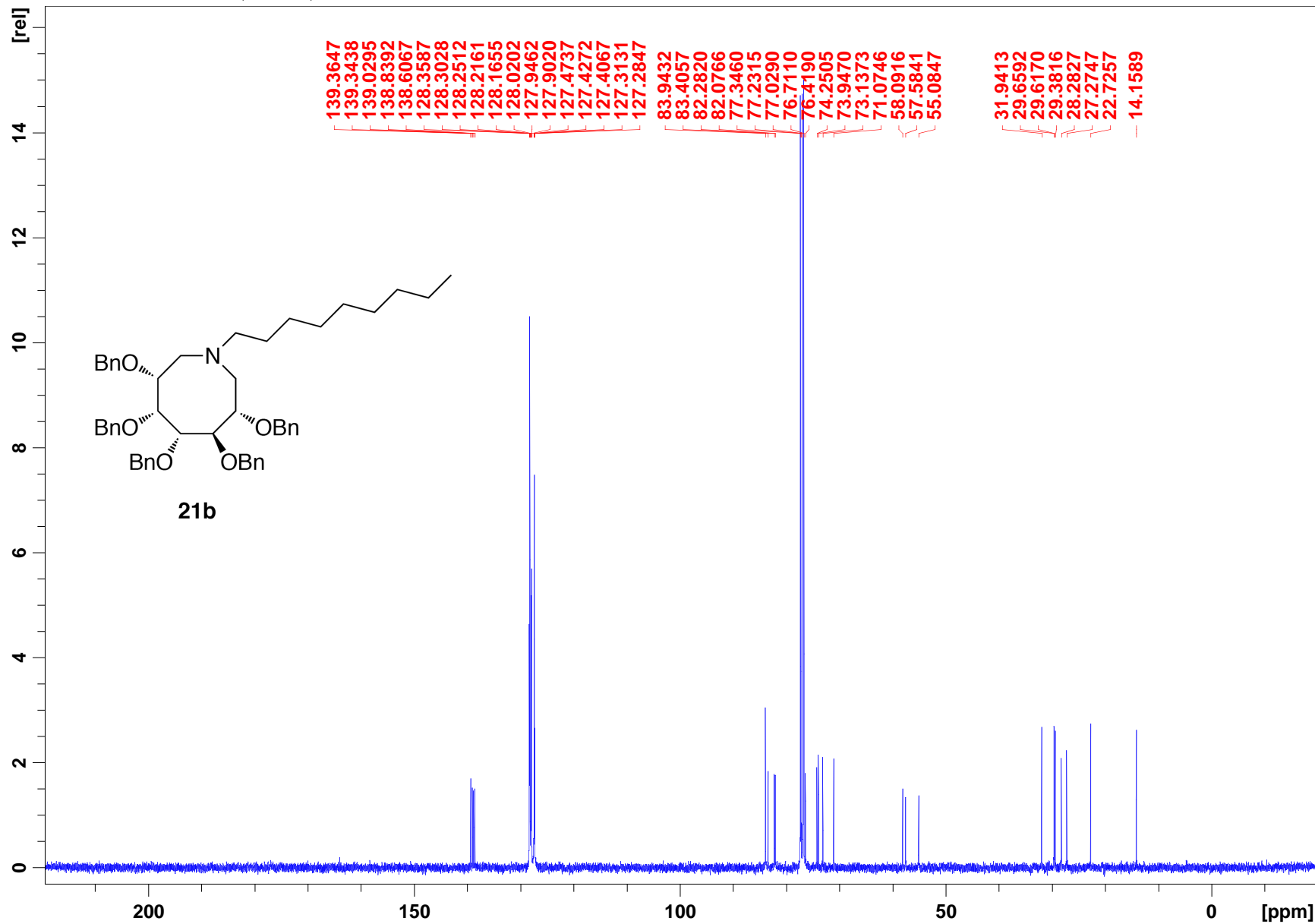
1H-NMR

JL-112207 1 1 C:\Bruker\TOPSPIN leex3451



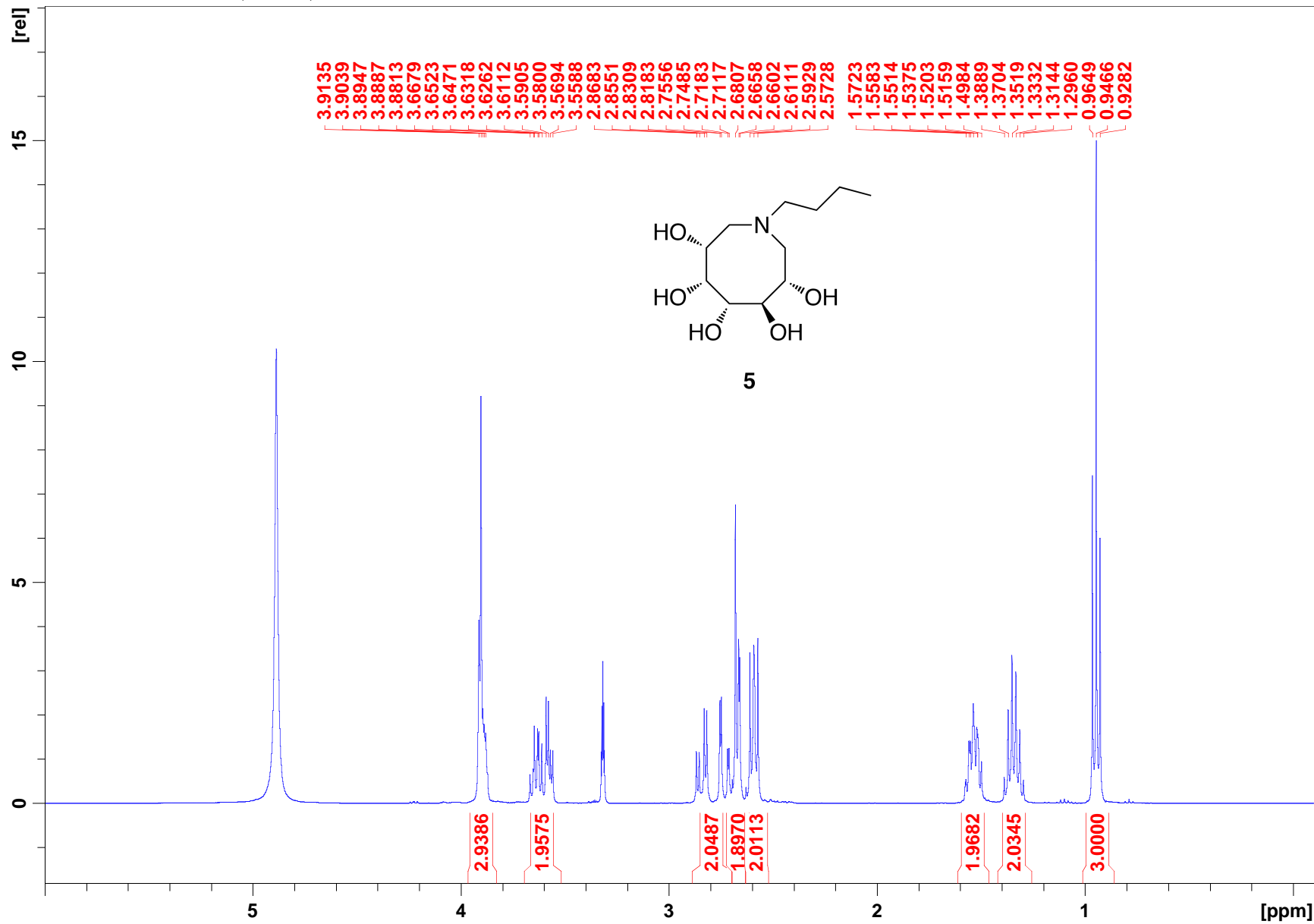
13C-NMR

JL-112207 3 1 C:\Bruker\TOPSPIN leex3451



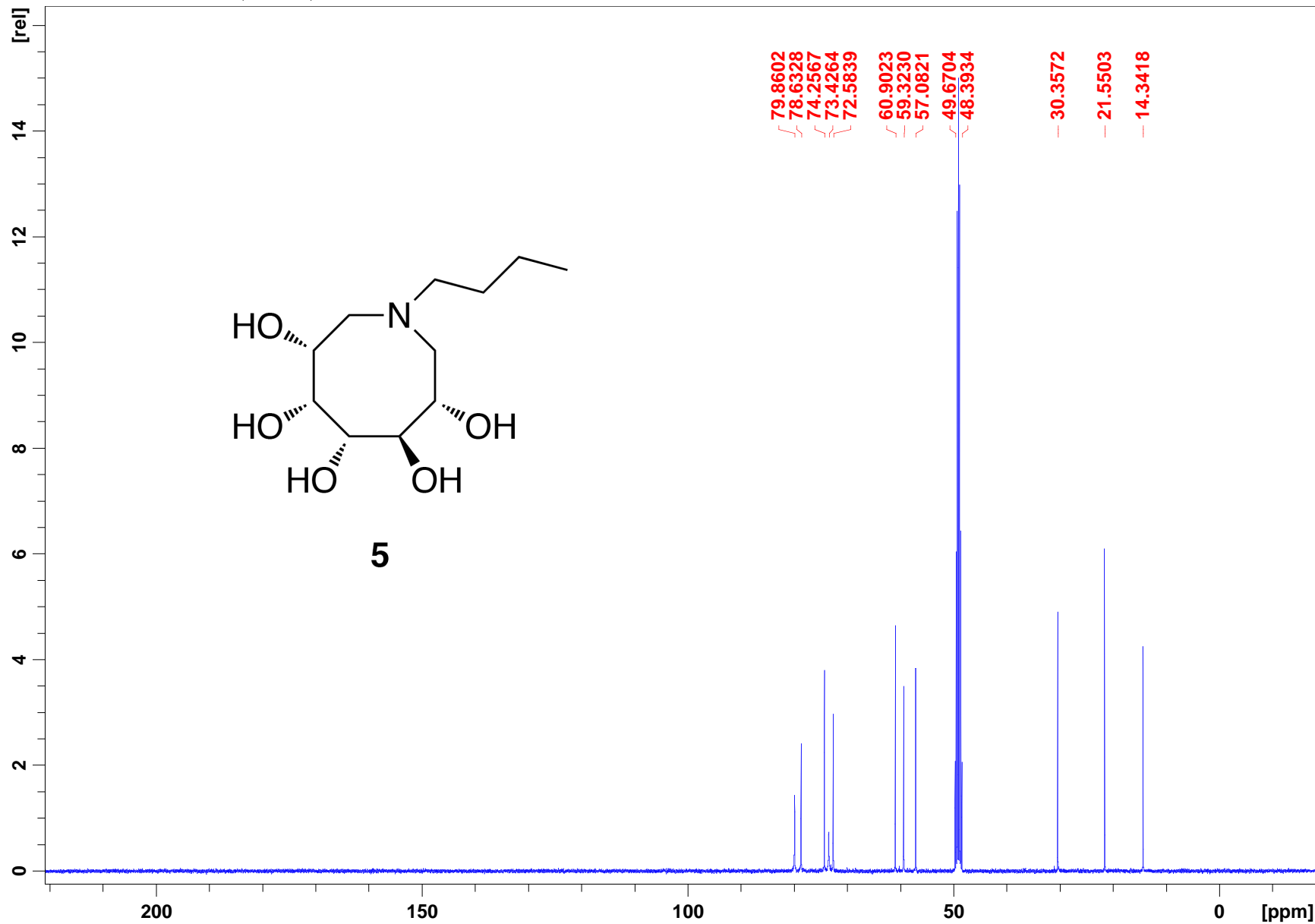
1H-NMR

JL-022008 1 1 C:\Bruker\TOPSPIN leex3451



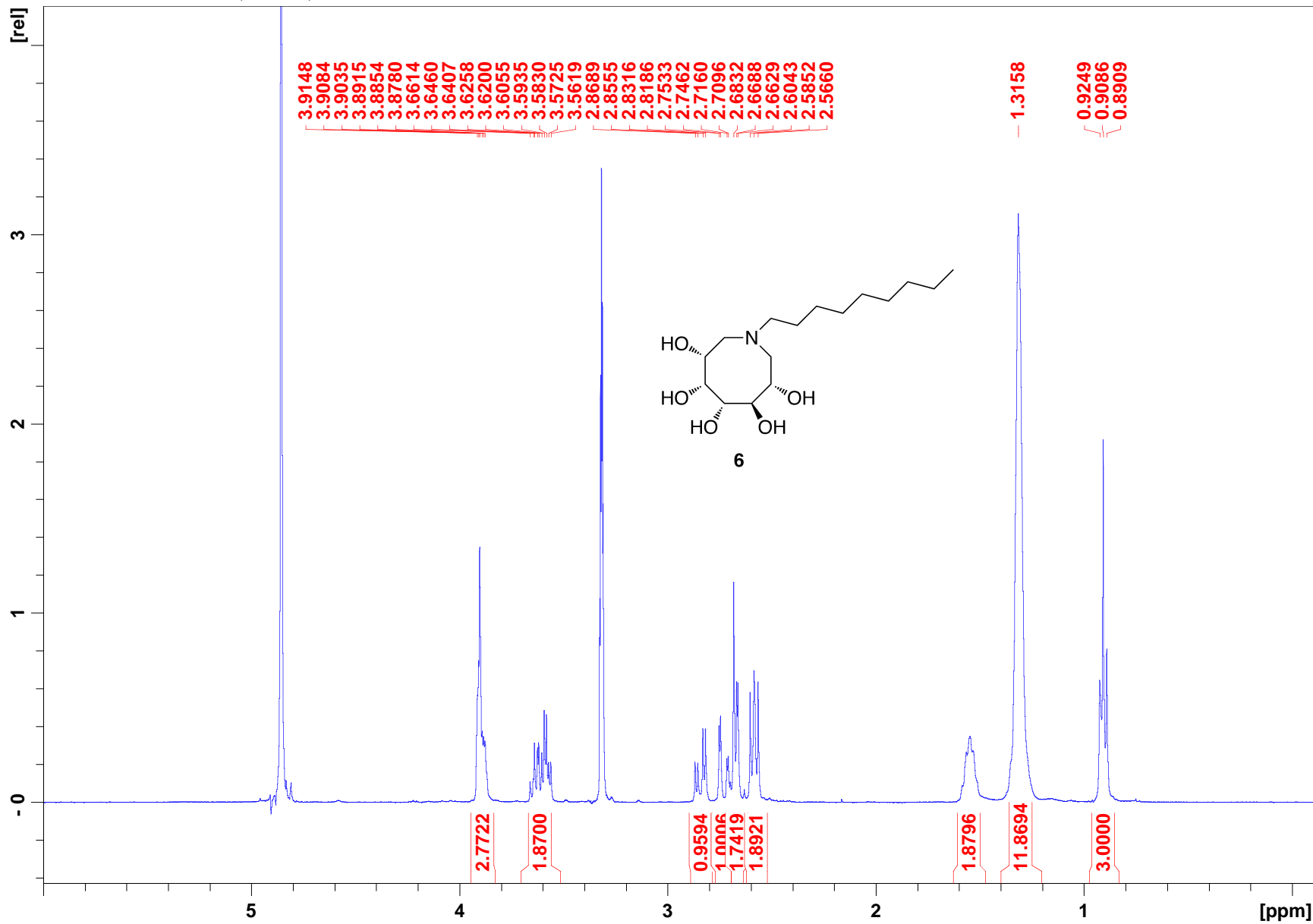
13C-NMR

JL-022008 3 1 C:\Bruker\TOPSPIN leex3451



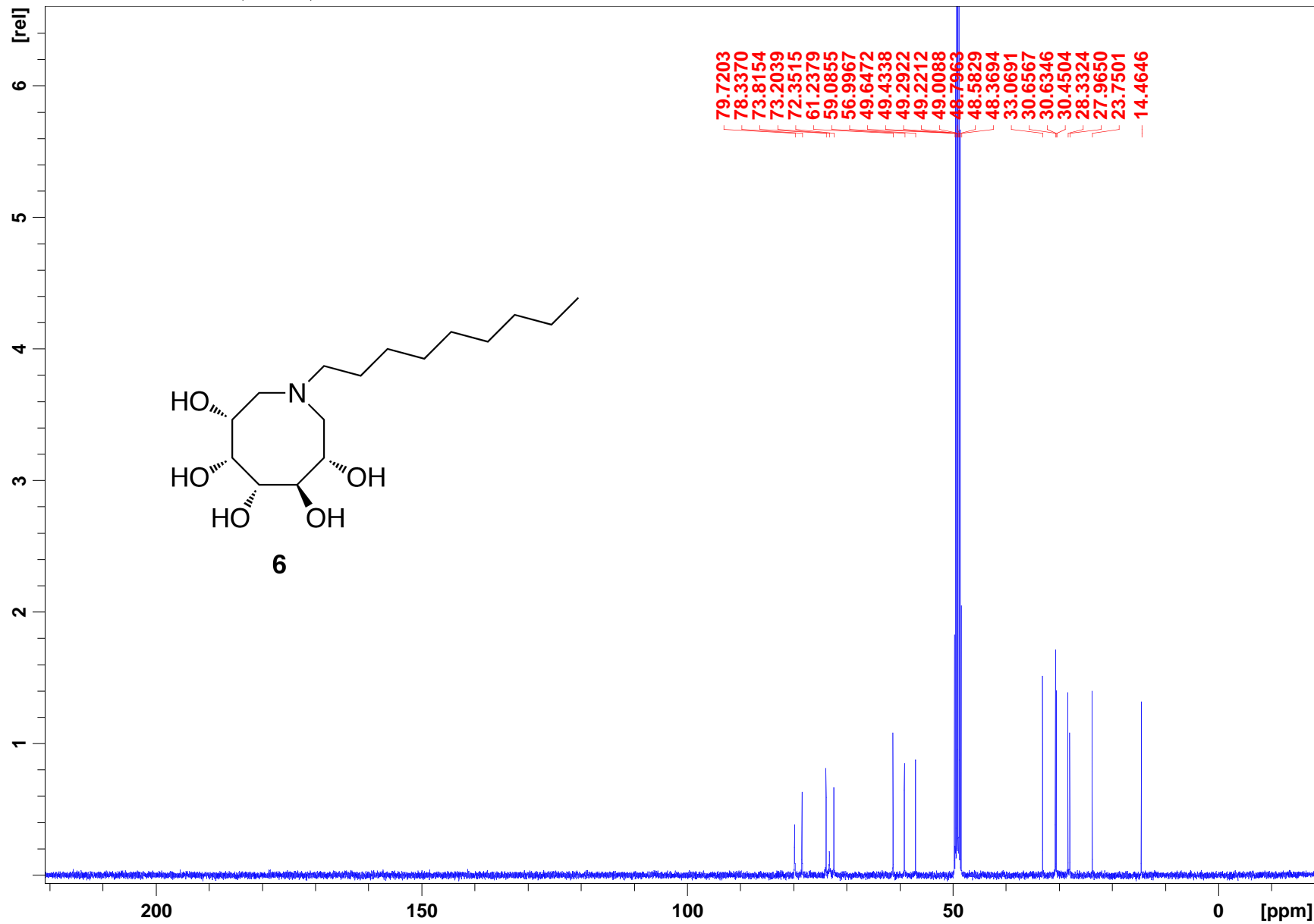
1H-NMR

JL-120507 1 1 C:\Bruker\TOPSPIN leex3451

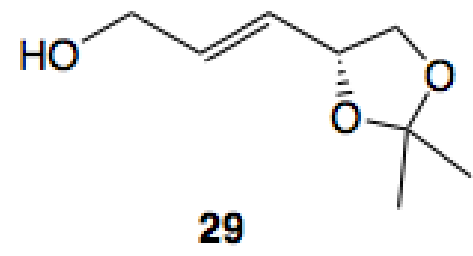


13C-NMR

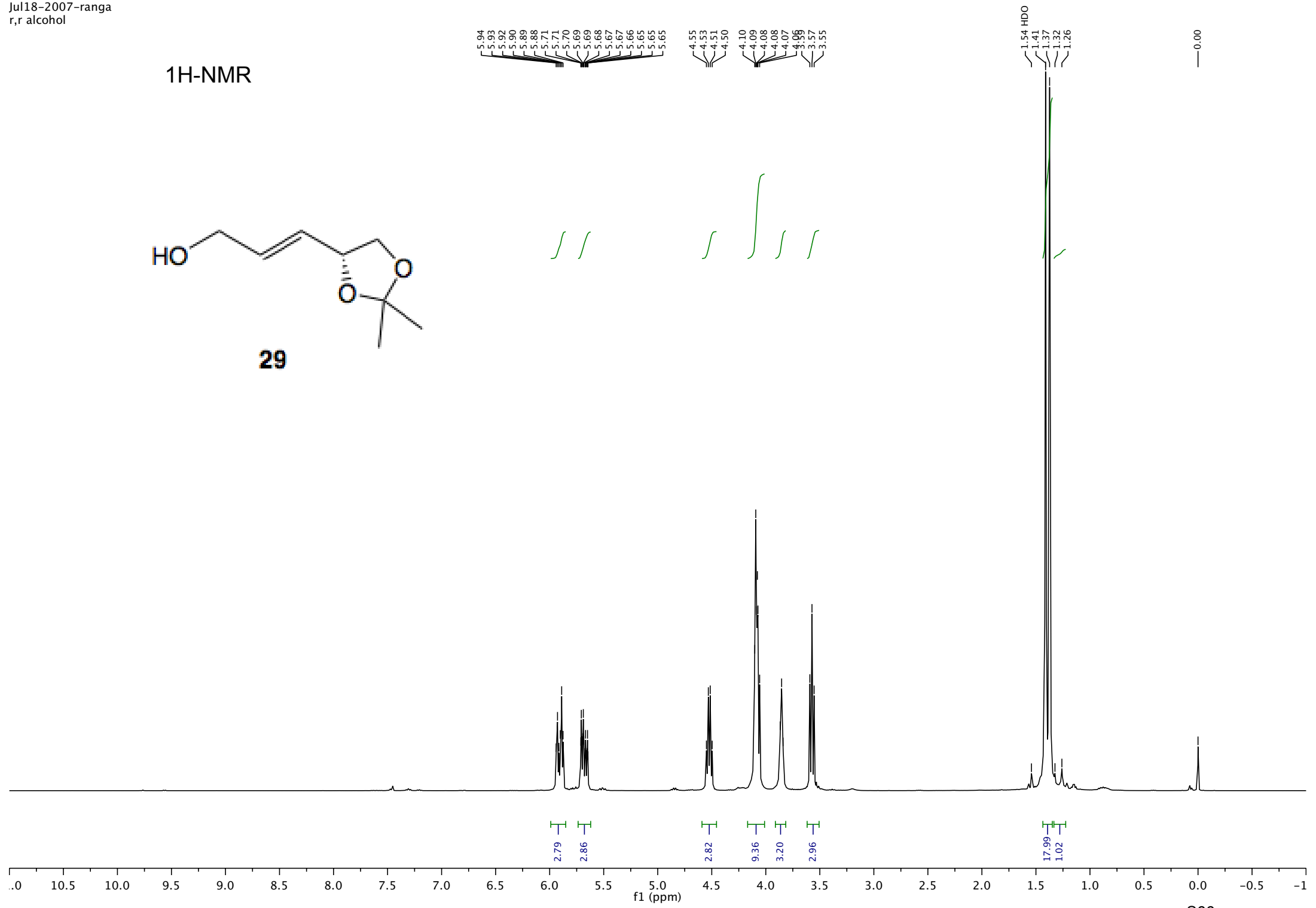
JL-030308 3 1 C:\Bruker\TOPSPIN leex3451



1H-NMR



5.94
5.93
5.92
5.90
5.88
5.88
5.71
5.71
5.70
5.69
5.68
5.67
5.66
5.65
5.65
4.55
4.53
4.51
4.50
4.10
4.09
4.08
4.08
4.07
4.06
3.57
3.55



2.79

2.86

2.82

9.36

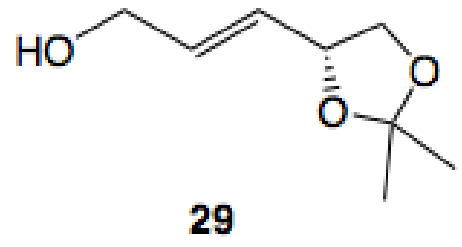
3.20

2.96

17.99

1.02

13C-NMR



133.71
133.41
128.88
127.81

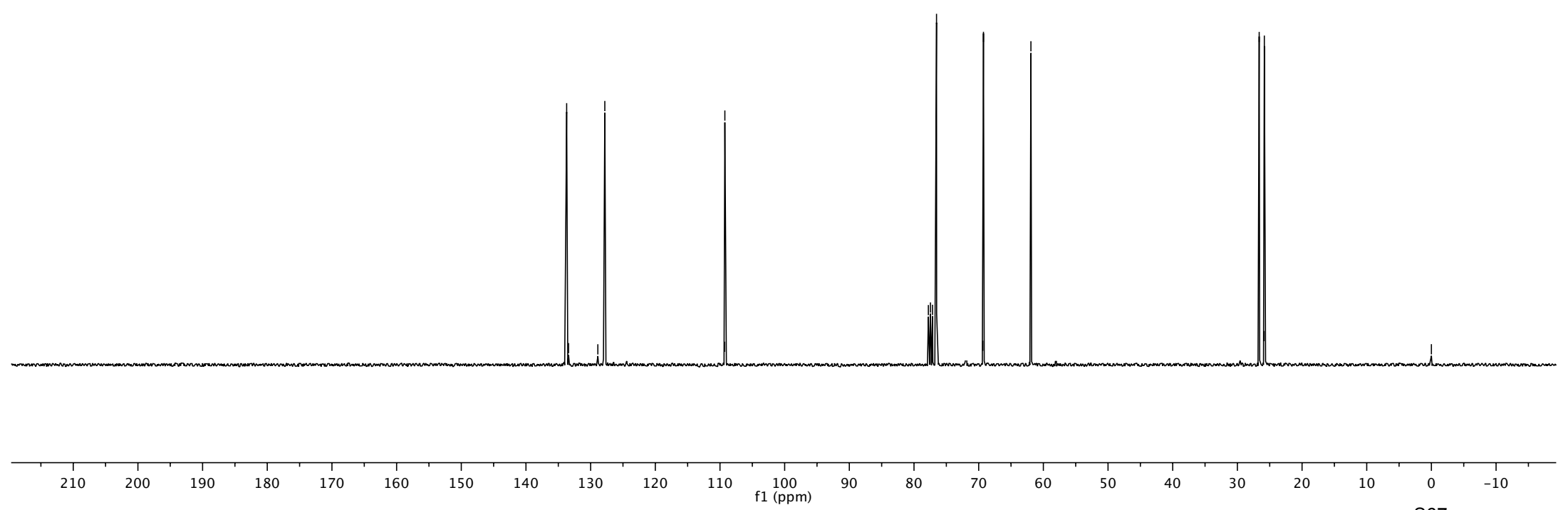
109.30
109.25

77.78
77.46
77.14
76.52 CDCl3
71.87
69.36
69.26

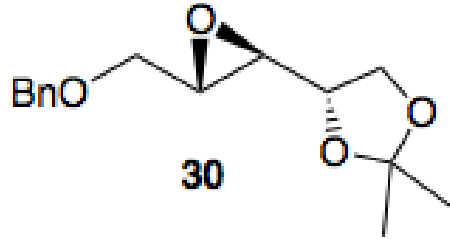
61.92
58.03

26.63
25.87
25.83

-0.00



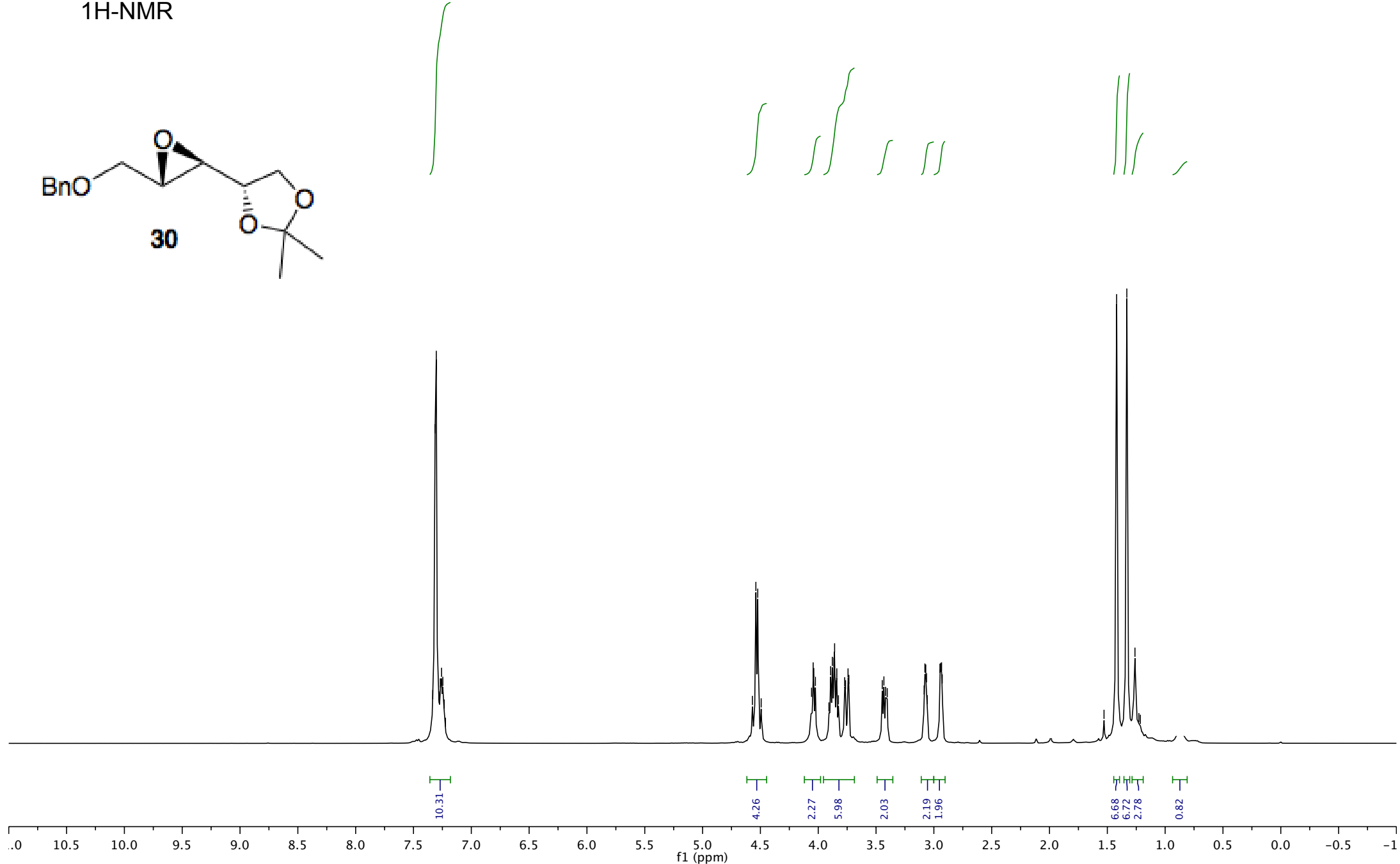
1H-NMR



7.33
7.31
7.30 CDCl3
7.28
7.27
7.26
7.25
7.23
7.22

4.57
4.54
4.52
4.49
4.04
3.89
3.88
3.86
3.84
3.45
3.43
3.42
3.08
3.07
3.06
2.95
2.94
2.93
2.93

— 1.99
1.53 H2O
1.42
1.33
1.26
1.23
1.21
0.90
0.88
0.85



13C-NMR

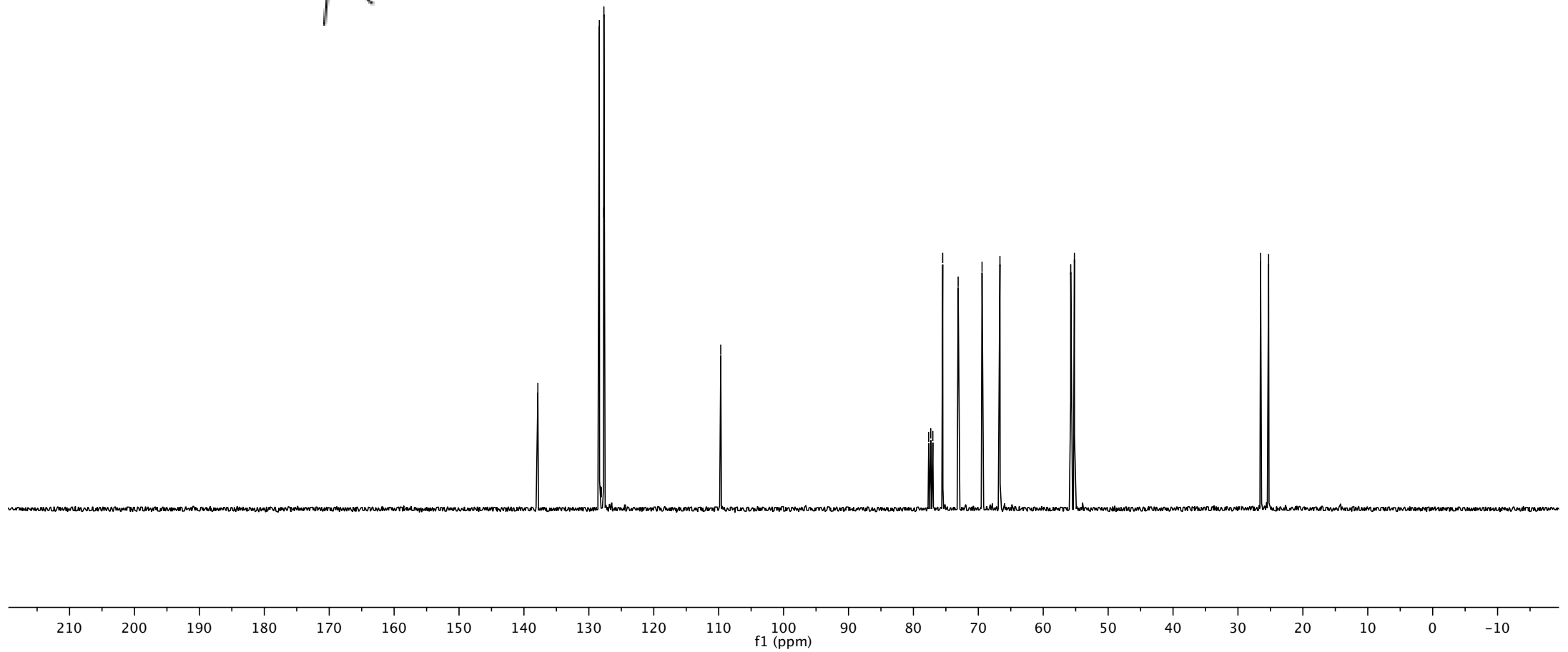
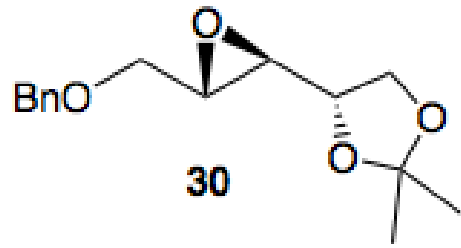
137.86
128.38
128.18
128.07
127.71
127.68
124.43

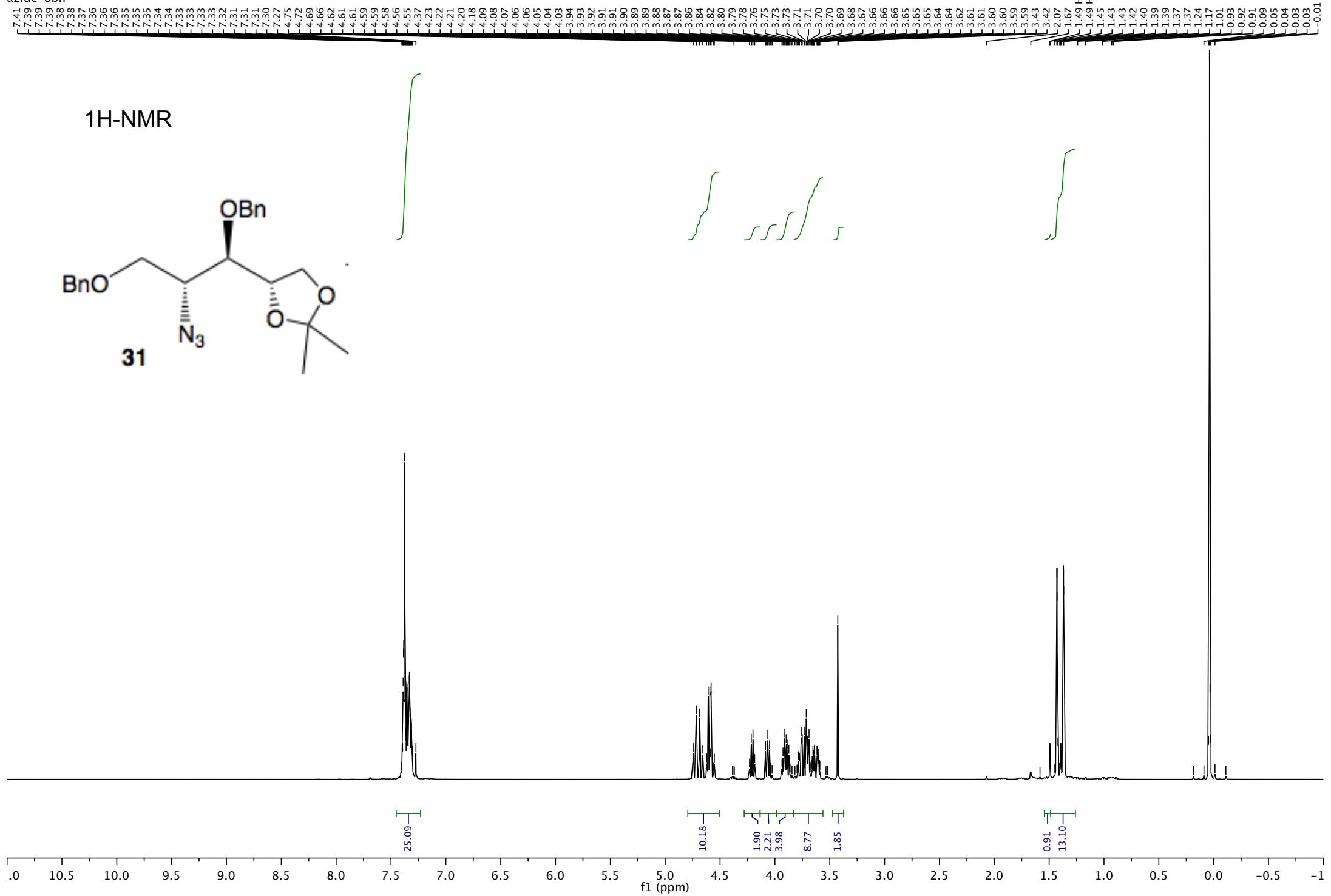
109.68

77.65
77.33
77.01 CDCl3
75.49
73.11
69.43
66.67

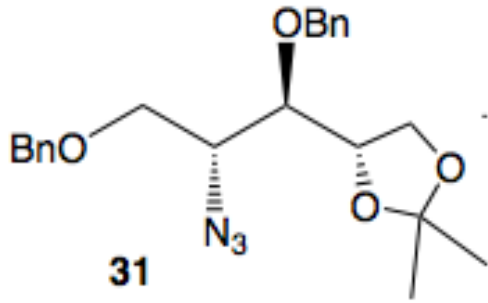
55.76
55.19

31.74
29.69
26.51
25.28





13C-NMR



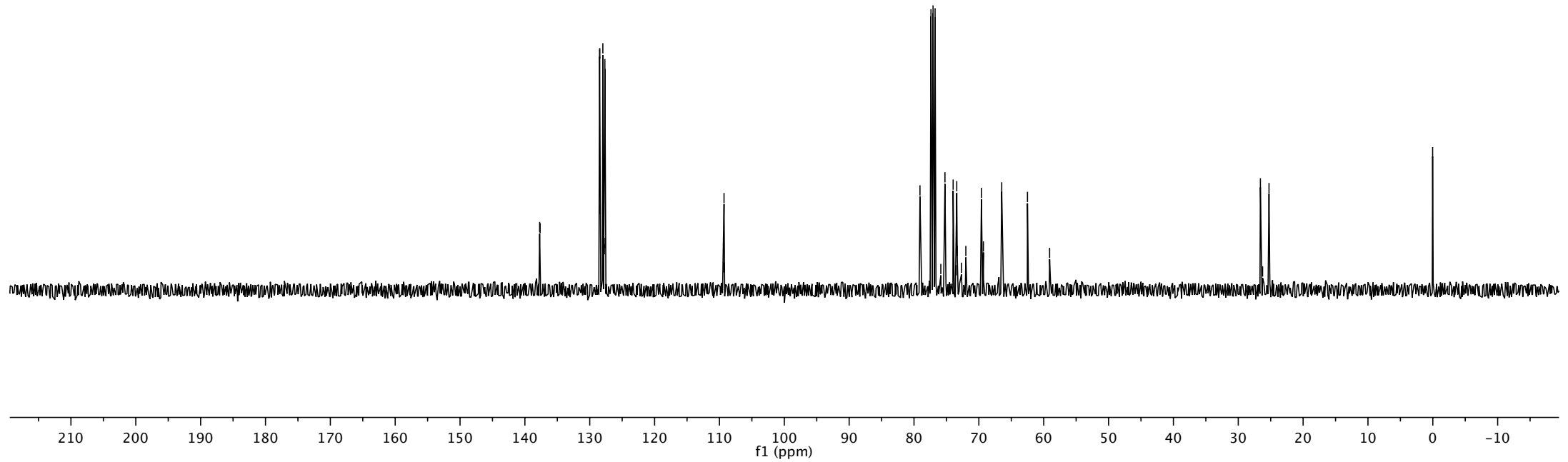
137.72
137.65
128.48
128.46
128.38
128.01
127.98
127.97
127.93
127.79
127.75
127.67
127.62
109.33
109.30

79.08
77.38 CDCl3
77.06 CDCl3
76.75 CDCl3
75.87
75.23
73.97
73.50
73.41
73.34
72.68
72.01
69.60
69.30
66.47
62.50
59.09

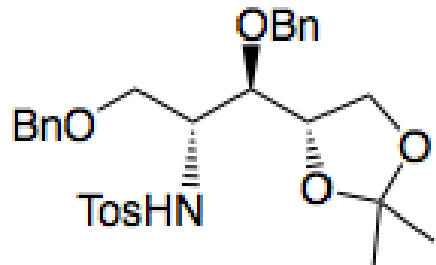
36.65

26.59
26.25
25.26

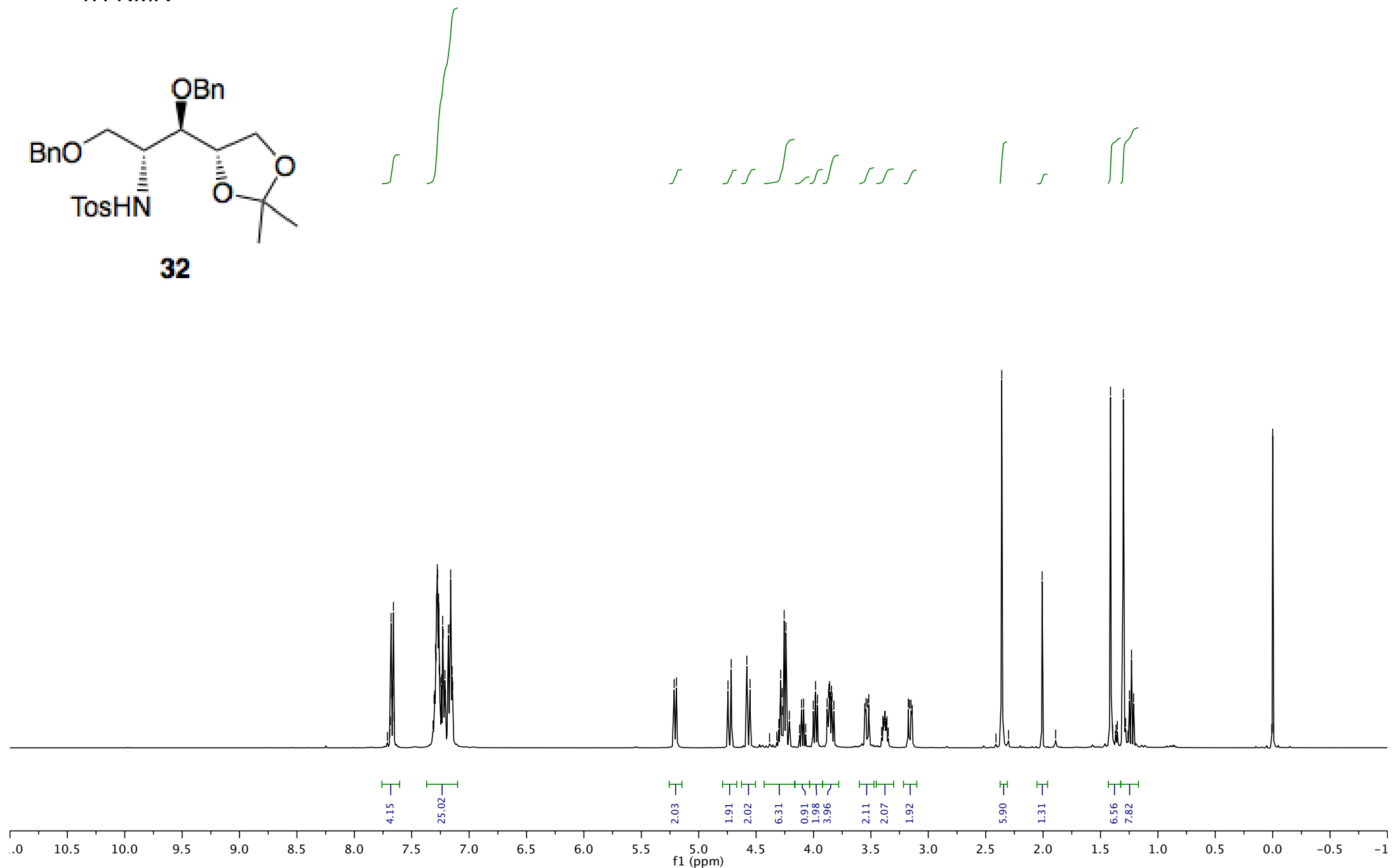
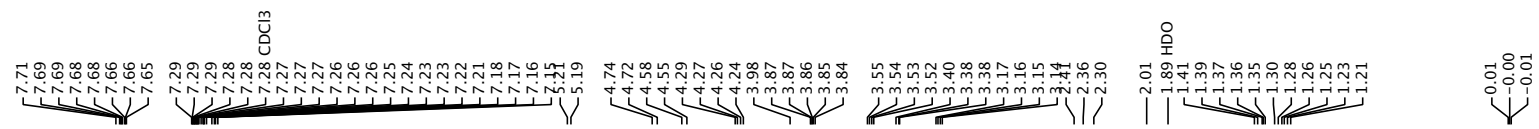
0.03



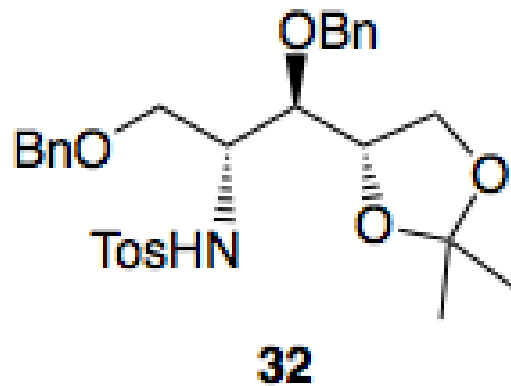
1H-NMR



32



13C-NMR



—171.04

—143.38
—138.35
—138.07
—137.49
—137.23
—129.57
—129.45
—128.34
—128.32
—128.29
—127.86
—127.83
—127.76
—127.70
—127.68
—127.13
—108.99

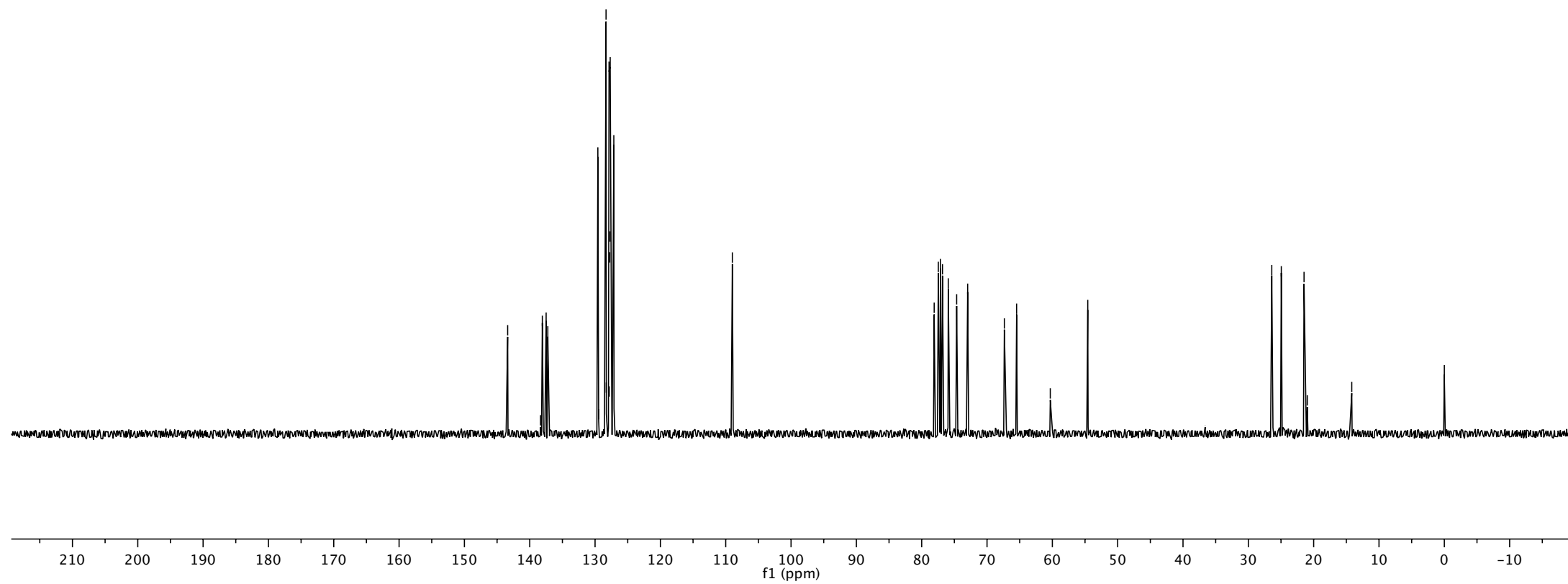
78.09
77.46
77.14 CDCl3
76.82
75.93
74.65
72.96
67.35
65.47
60.32
54.58

—36.62

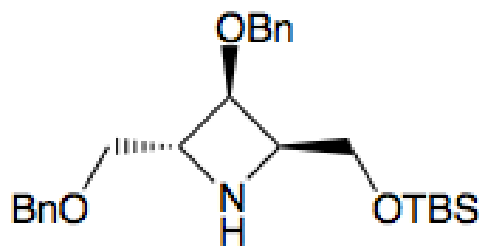
—26.41
—24.95
—21.47
—20.98

—14.17

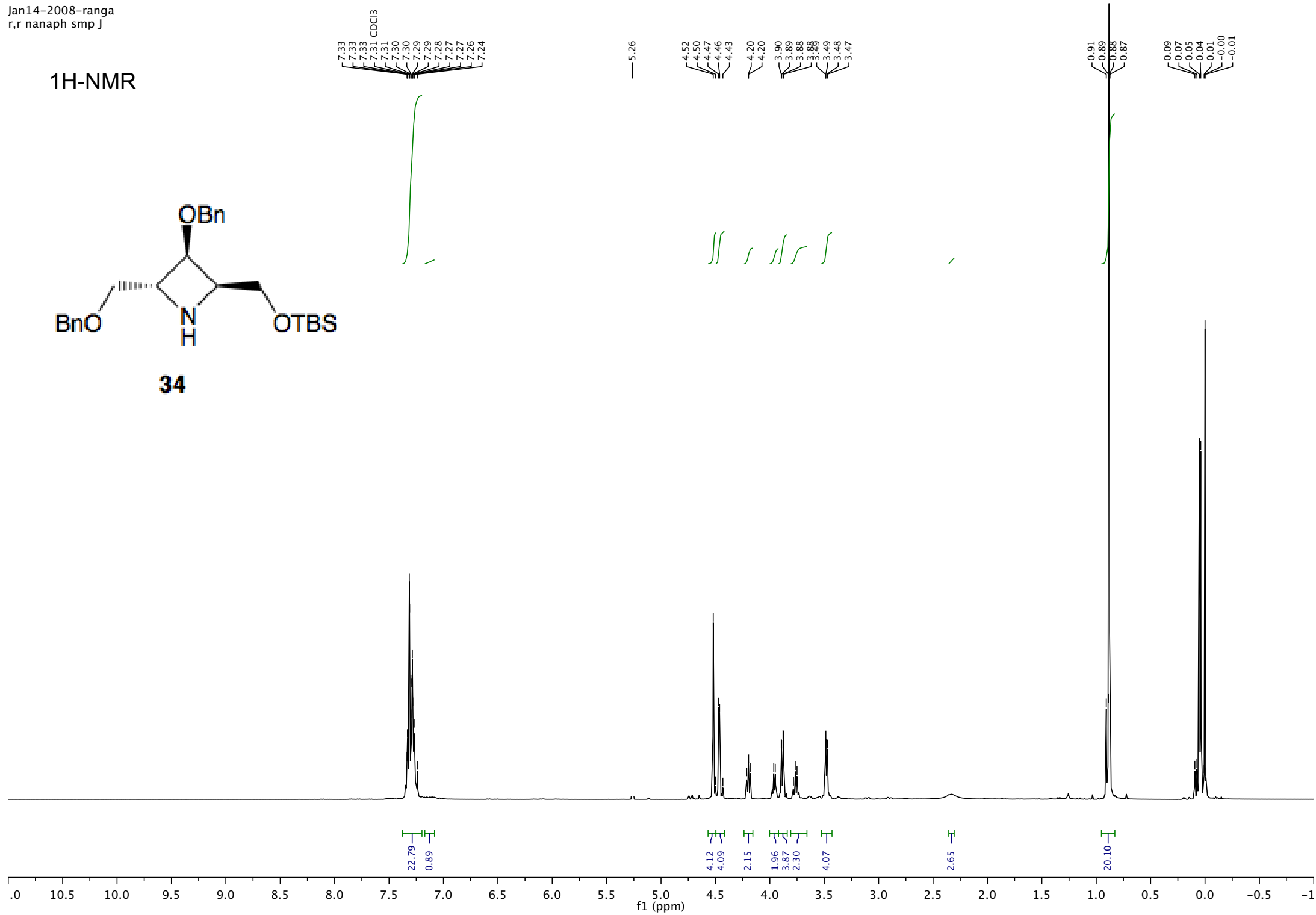
—0.00



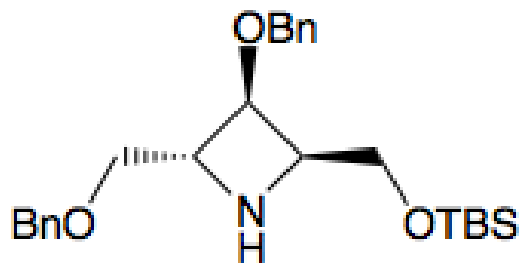
1H-NMR



34



13C-NMR

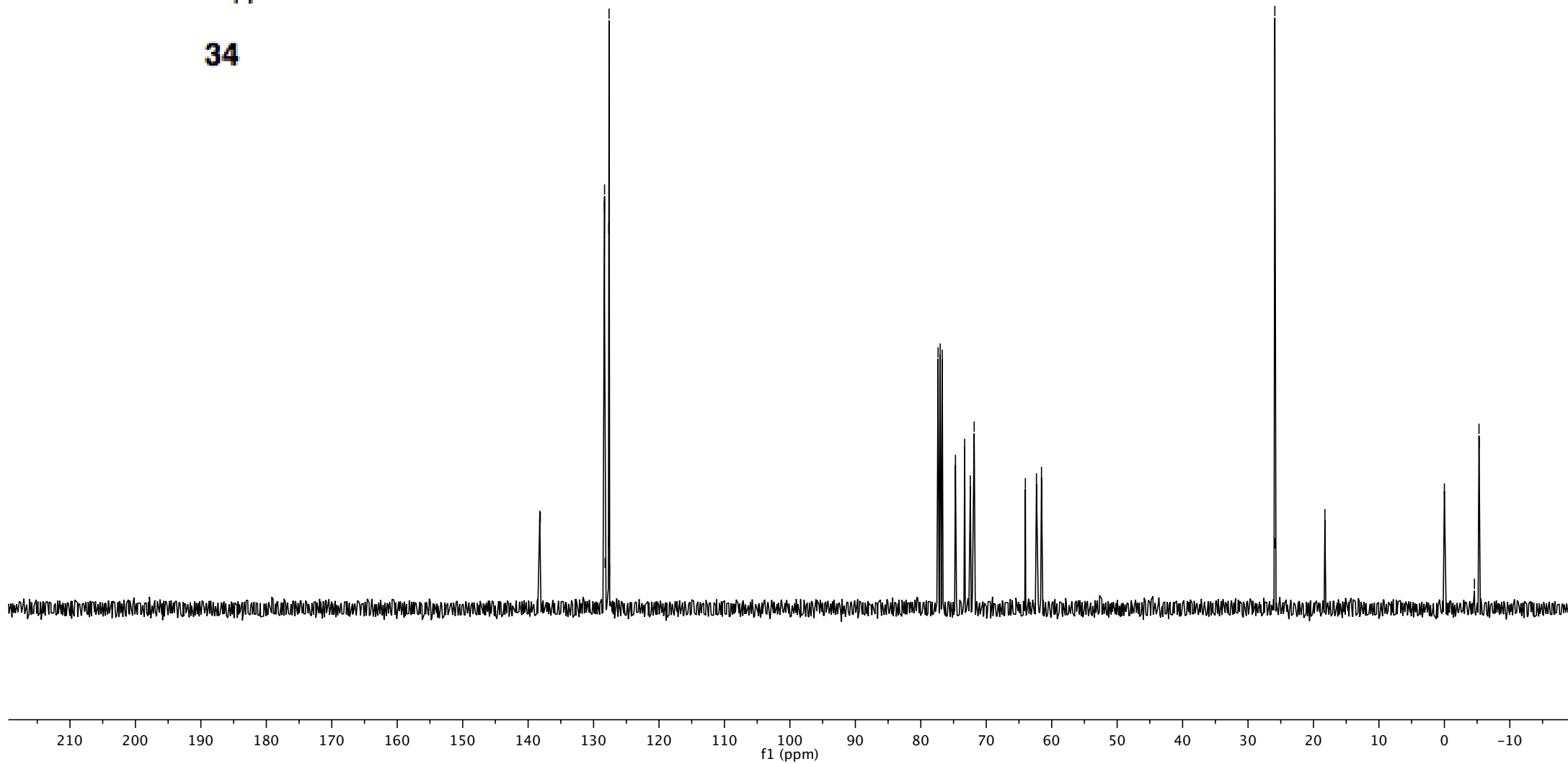


138.23
138.14
128.35
128.32
128.24
127.72
127.68
127.63
127.59
127.56

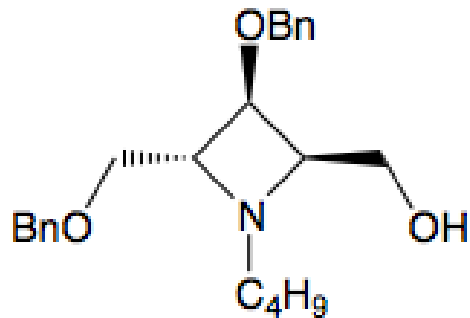
80.59
77.36
77.04 CDCl3
76.73 CDCl3
74.71
73.30
72.43
71.84
64.03
62.33
61.56

25.91
25.86
18.26

0.00
-4.59
-5.29
-5.31



1H-NMR



35a

7.36
7.36
7.35
7.34
7.34
7.33
7.33
7.32
7.31
7.30
7.30
7.29
7.29
7.28
7.28
7.28
7.27
7.27
7.26
7.26
7.25
7.25

CDCl₃

5.28

4.64
4.56
4.55
4.53
4.52
4.49
4.46
4.43

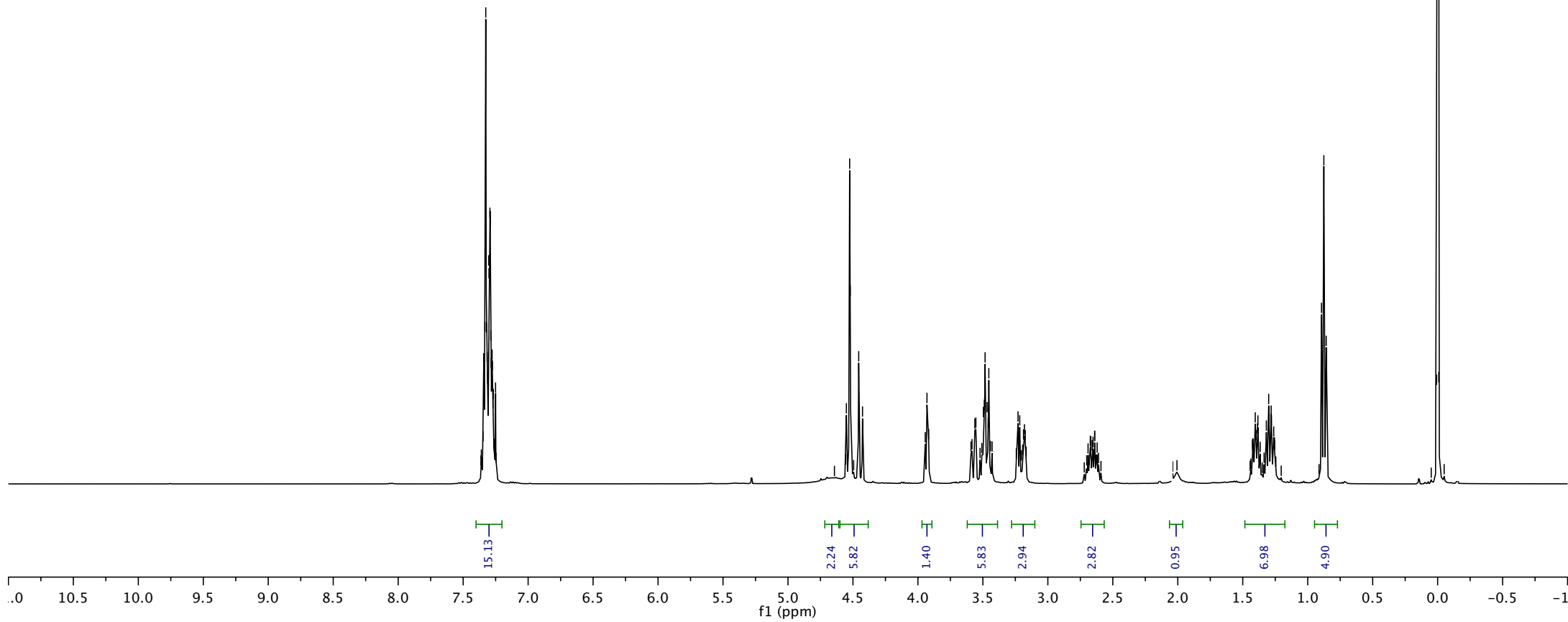
3.94
3.93
3.92

3.55
3.50
3.49
3.48
3.47
3.45
3.23
3.22

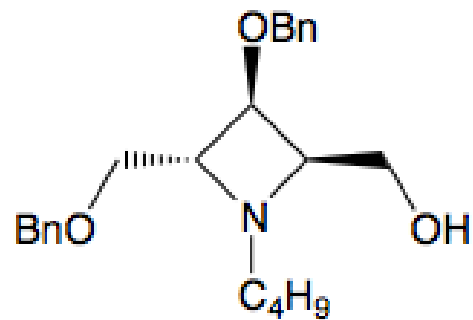
2.67
2.66
2.65
2.64
2.64
2.64
2.64
2.04
2.01

1.40
1.38
1.32
1.30
1.28
1.28
0.89
0.88
0.86

0.15
0.05
0.01
0.01
-0.01
-0.05
-0.15



13C-NMR



35a

138.16
137.91
128.44
128.41
128.36
127.84
127.61
127.57

77.34 CDCl₃
77.03 CDCl₃
76.71 CDCl₃
73.35
72.35
71.64
71.63
71.25
70.09

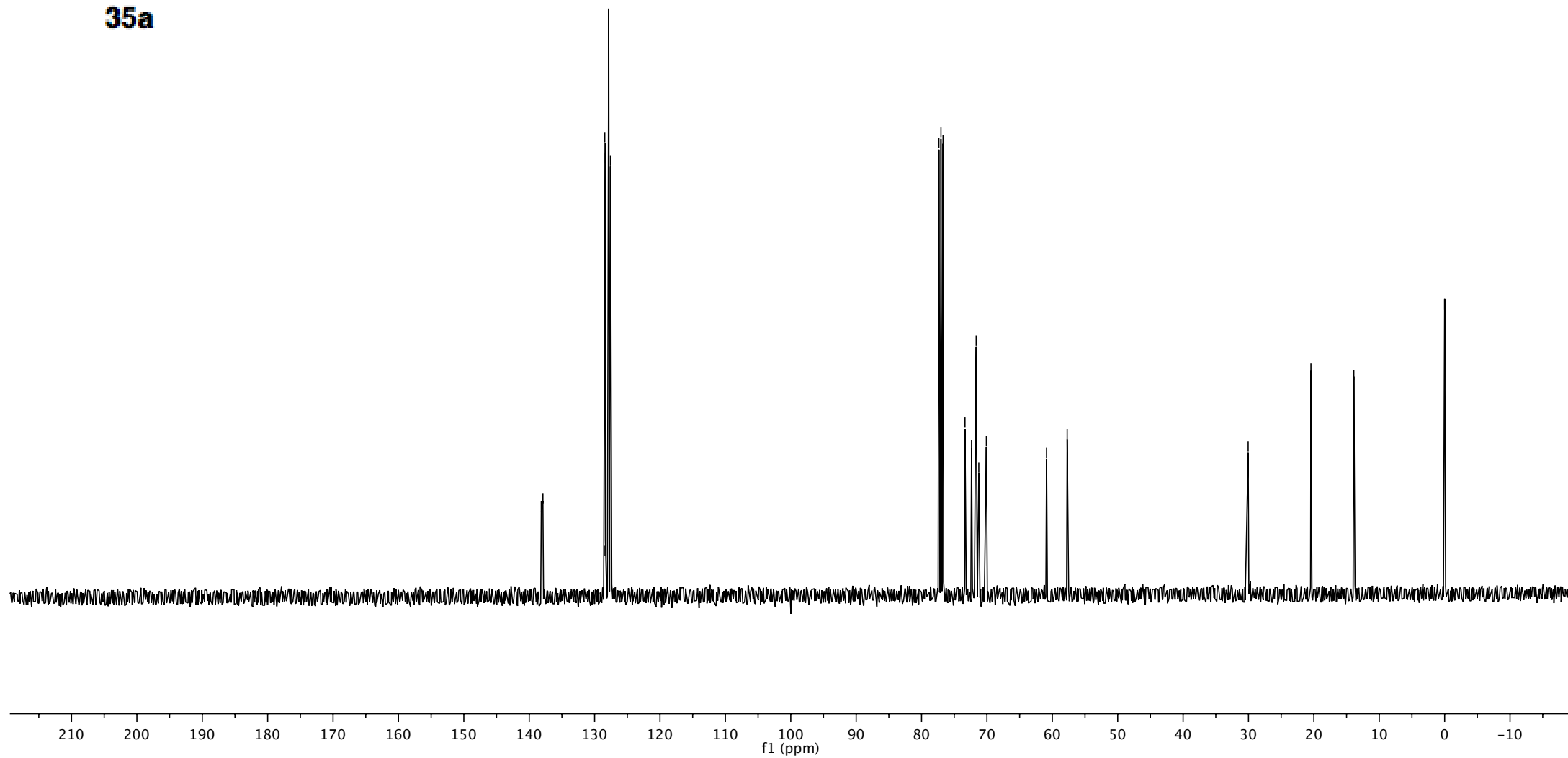
60.88
57.73

30.05

20.45

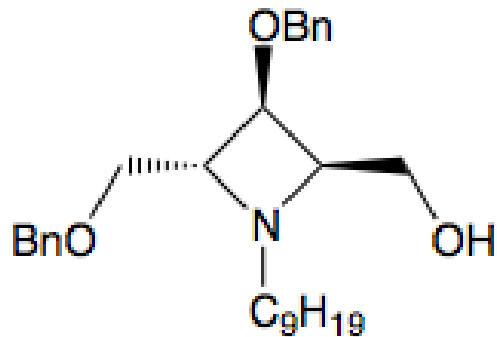
13.89

0.00

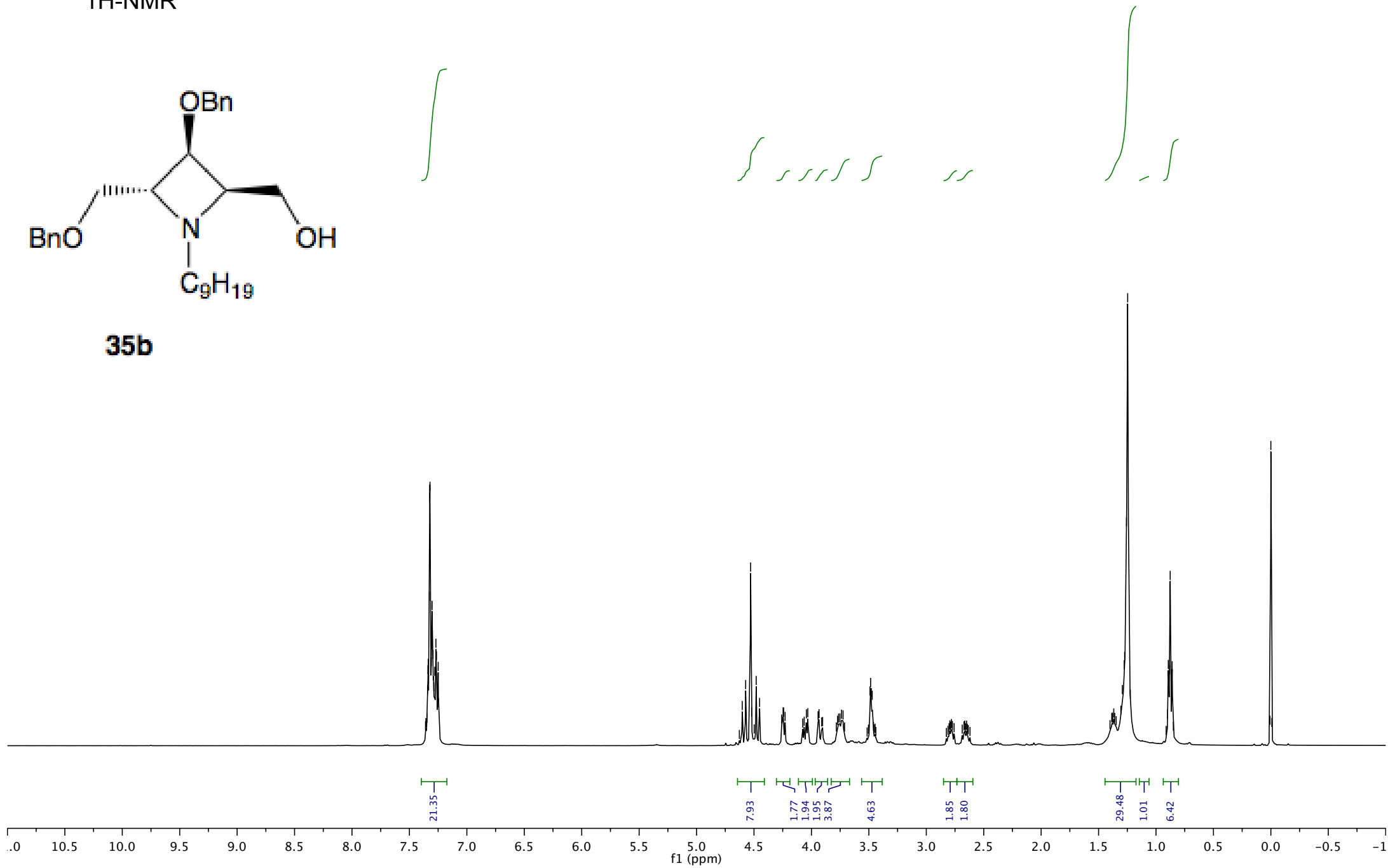


7.36
7.36
7.35
7.34
7.34
7.33
7.32
7.31
7.31
7.30
7.29
7.28
7.27
7.26
7.26 CDCl3
7.25
7.25
4.60
4.57
4.53
4.52
4.48
4.45
4.26
4.25
4.24
4.23
4.08
4.06
4.04
4.03
3.94
3.93
3.91
3.90
3.78
3.77
3.76
3.75
3.74
3.72
3.71
3.49
3.48
3.48
3.47
3.46
3.45
2.80
2.79
2.79
2.78
2.77
2.76
2.67
2.66
2.66
2.65
2.64
1.38 H2O
1.37 H2O
1.31
1.29
1.28
1.27
1.26
1.25
0.91
0.89
0.88
0.88
0.86
0.85
0.01
0.00
-0.01

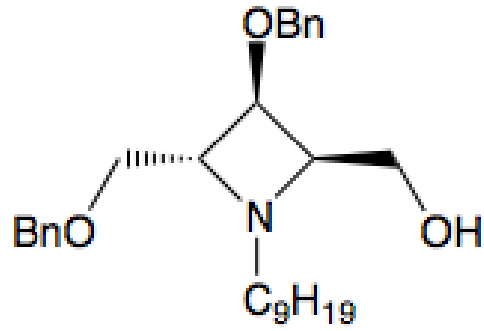
1H-NMR



35b



13C-NMR



35b

138.14
137.58
128.49
128.44
128.37
127.95
127.90
127.71
127.69
127.65
127.61

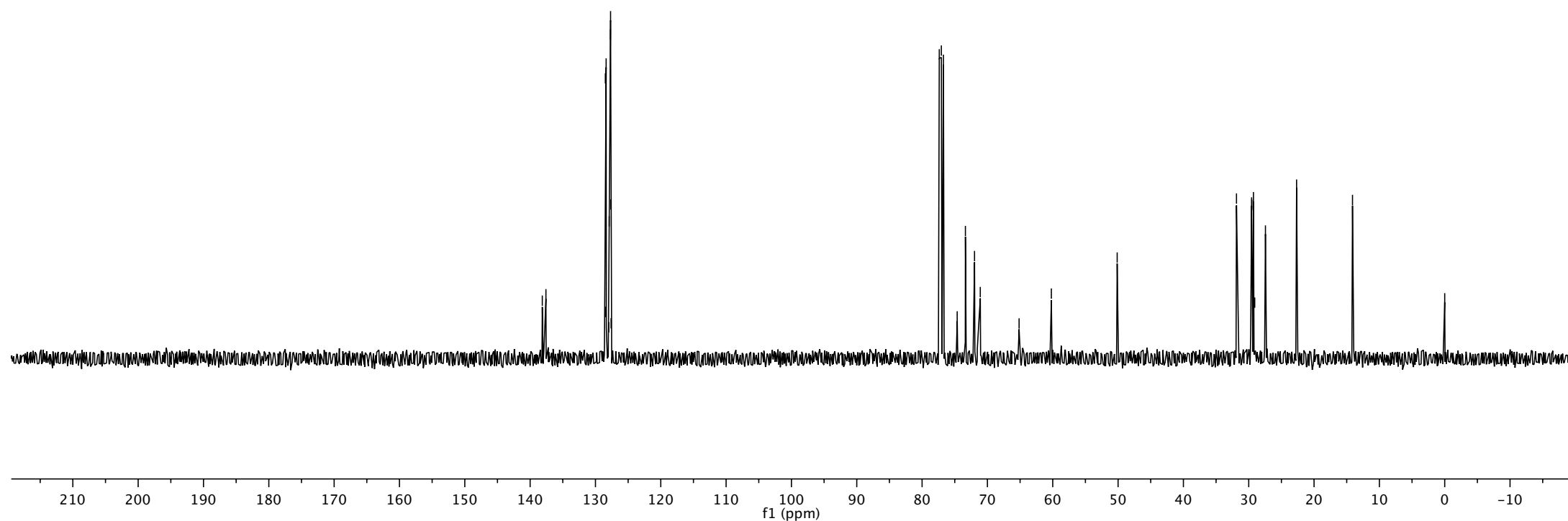
77.37 CDCl₃
77.05 CDCl₃
76.73
74.63
73.37
71.97
65.15
60.21

50.14

31.87
29.56
29.53
29.28
29.08
27.44
22.67

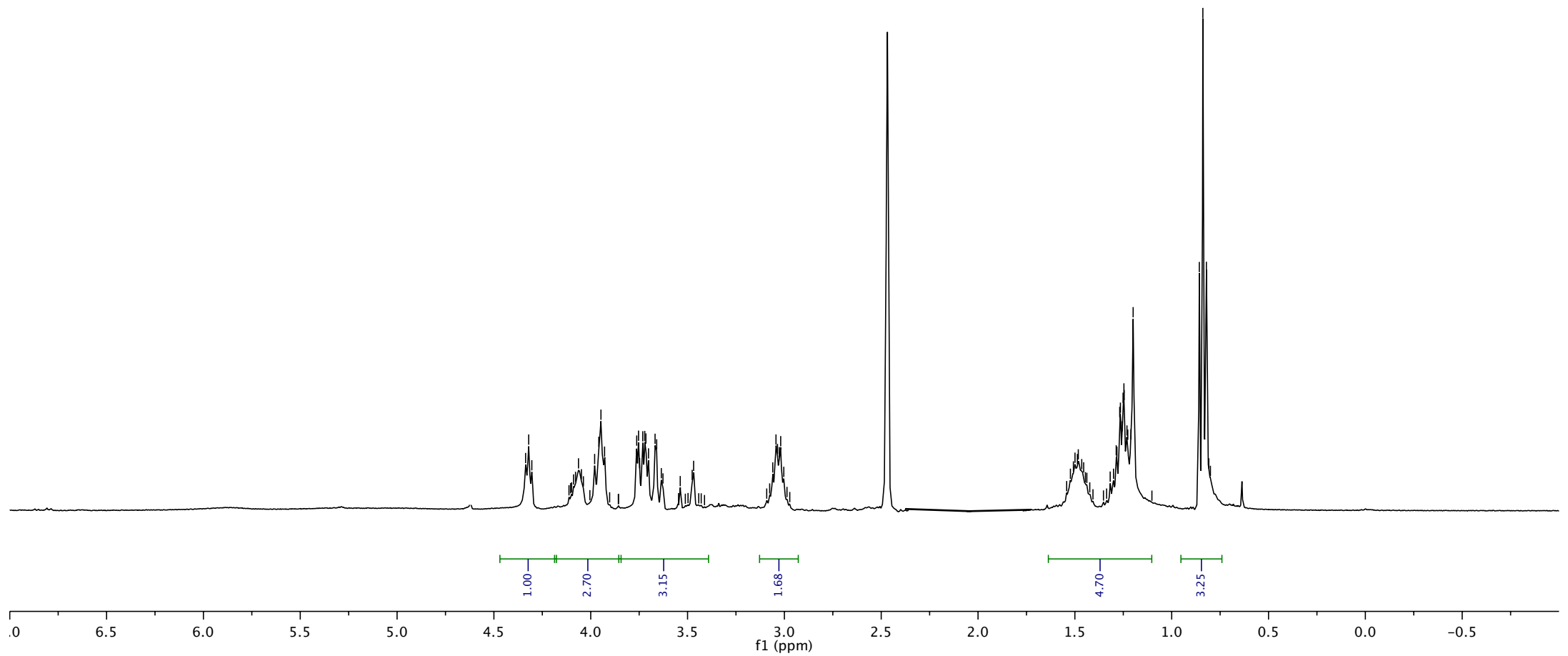
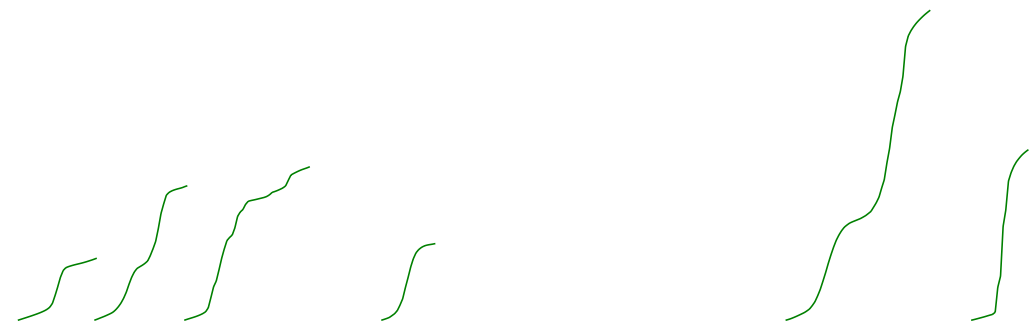
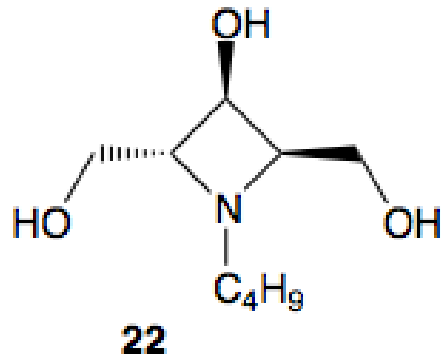
14.12

-0.00

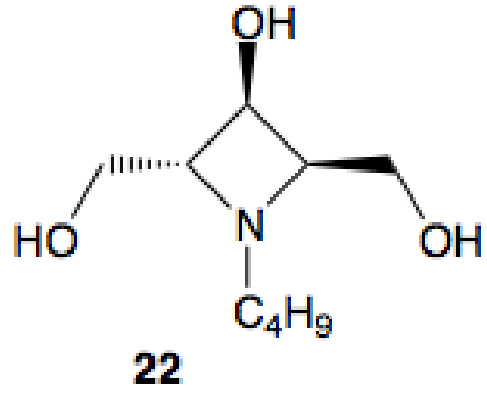


4.34
4.32
4.30
4.06
3.98
3.96
3.95
3.93
3.76
3.75
3.75
3.72
3.72
3.70
3.67
3.69
3.08
3.06
3.04
3.03
3.02
3.00
2.99
2.97
1.54
1.52
1.51
1.50
1.49
1.48
1.46
1.45
1.45
1.44
1.32
1.30
1.29
1.28
1.27
1.26
1.25
1.25
1.23
1.23
1.21
1.21
0.84
0.82
0.81
0.80

1H-NMR

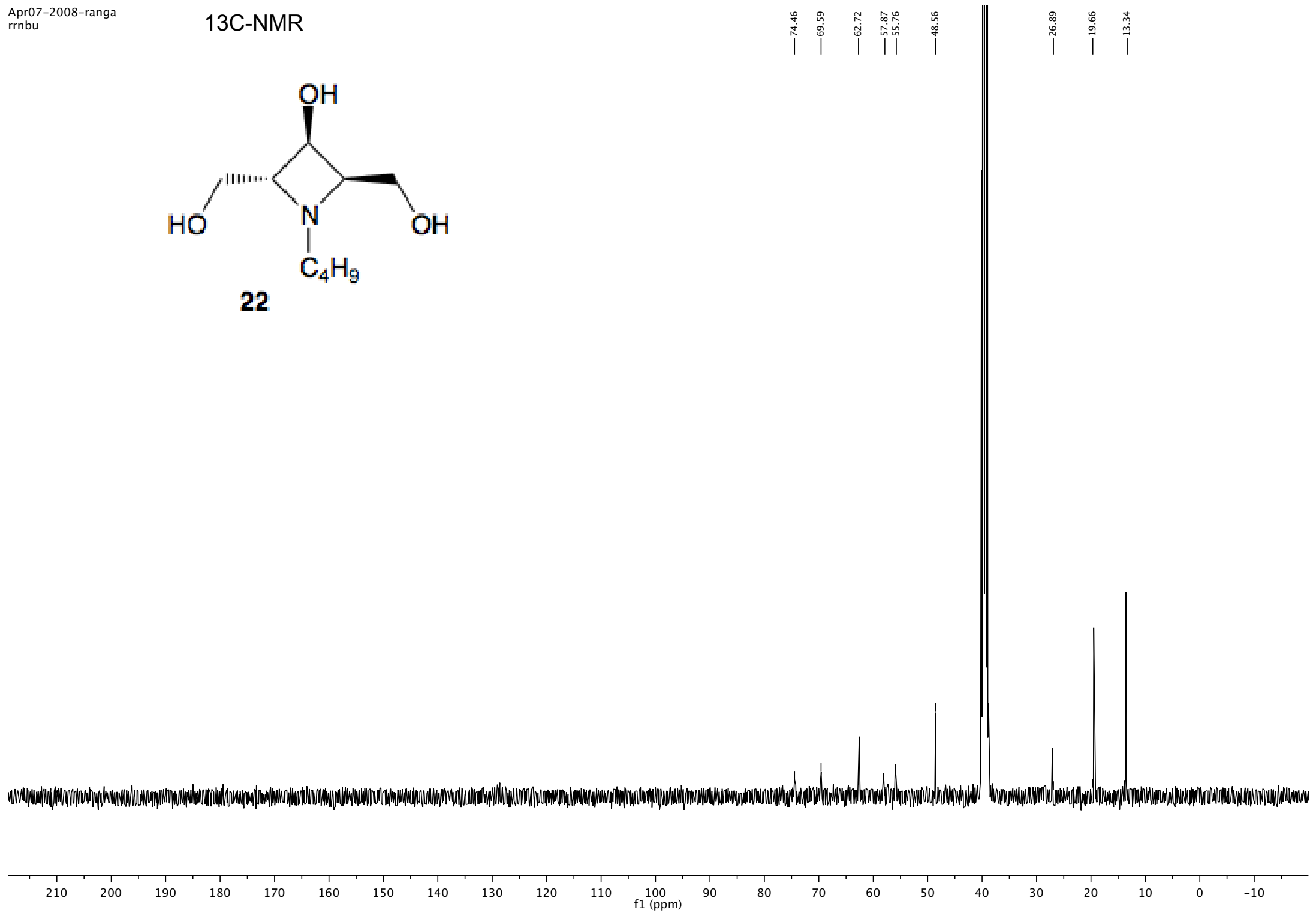


13C-NMR

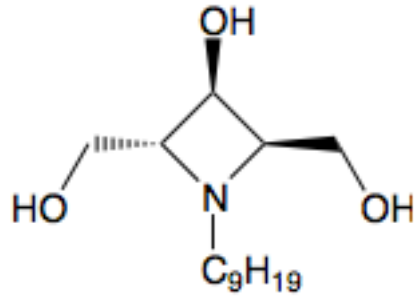


— 74.46
— 69.59
— 62.72
— 57.87
— 55.76
— 48.56

— 26.89
— 19.66
— 13.34

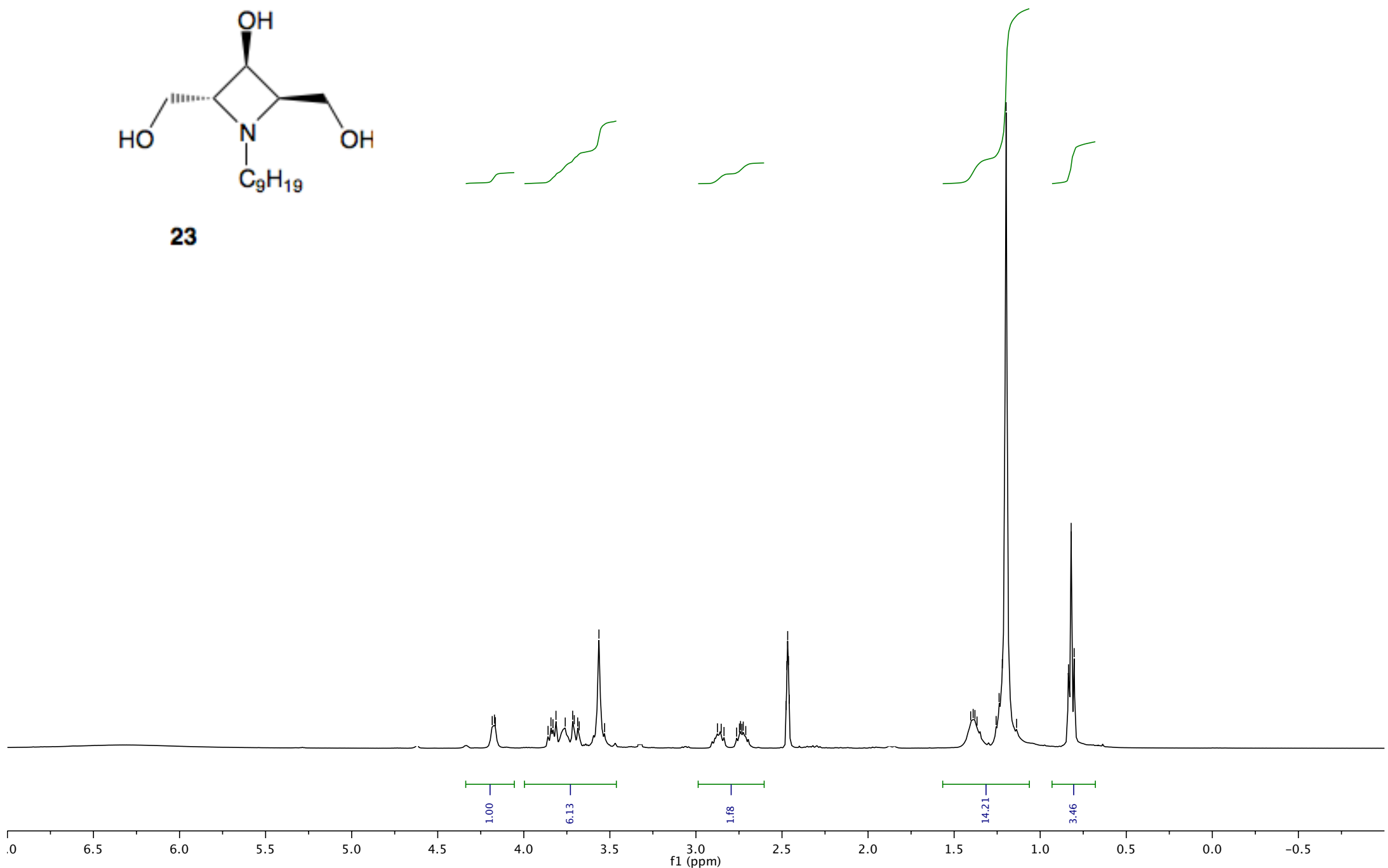


1H-NMR

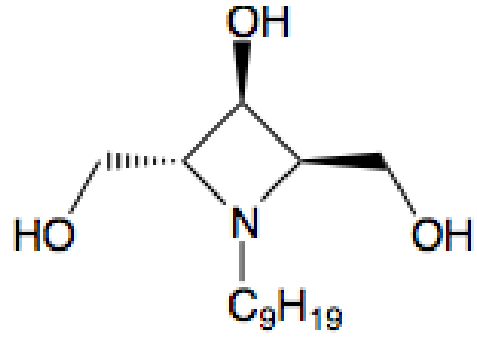


23

- 4.18
- 4.17
- 4.16
- 3.86
- 3.84
- 3.83
- 3.81
- 3.76
- 3.72
- 3.71
- 3.69
- 3.68
- 3.56 HDO
- 3.53 HDO
- 2.87
- 2.85
- 2.84
- 2.76
- 2.75
- 2.74
- 2.73
- 2.73
- 2.71
- 2.48 DMSO
- 2.47 DMSO
- 2.47 DMSO
- 2.46 DMSO
- 2.46 DMSO
- 1.40
- 1.39
- 1.38
- 1.37
- 1.26
- 1.24
- 1.22
- 1.20
- 1.14
- 0.84
- 0.82
- 0.80

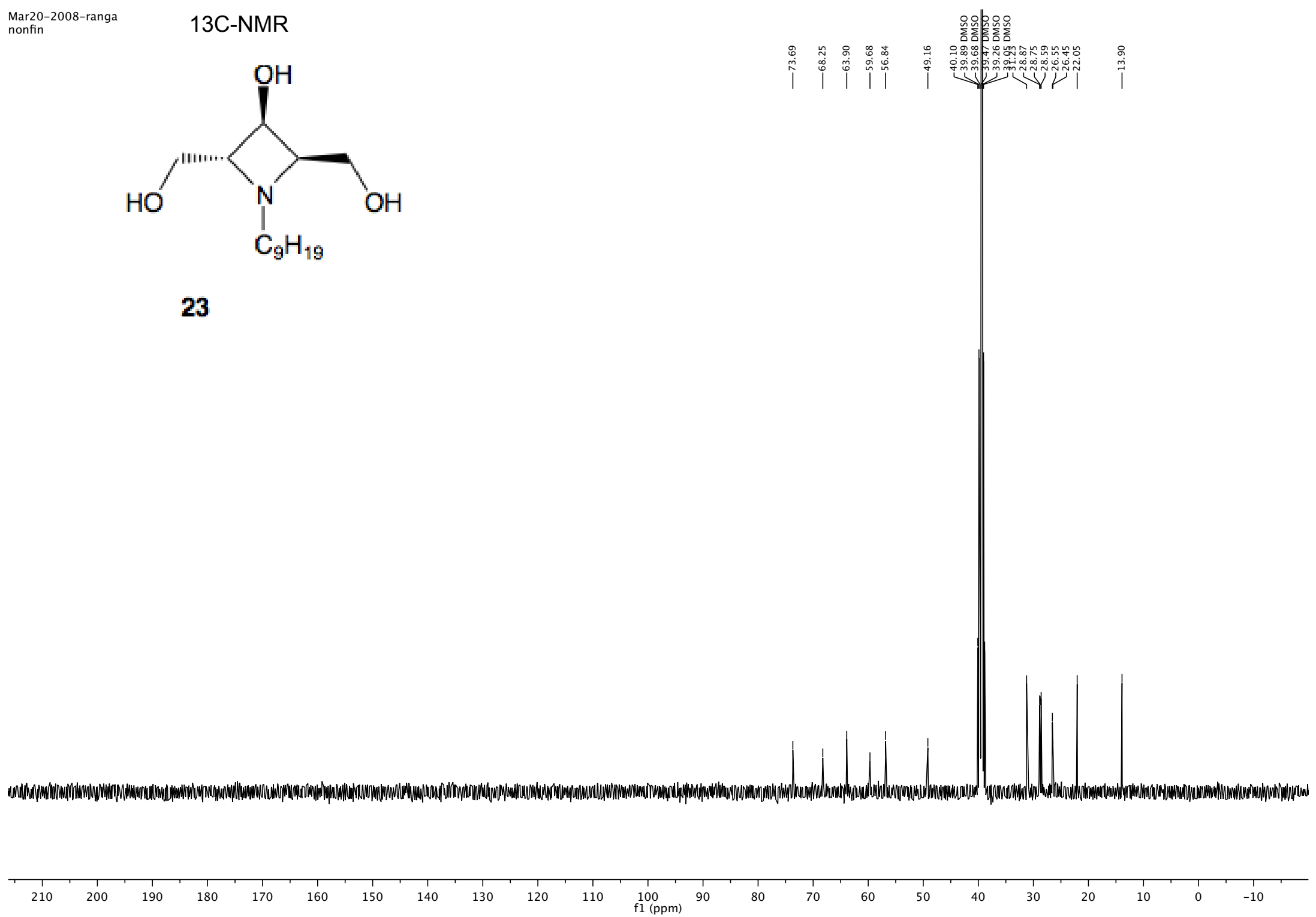


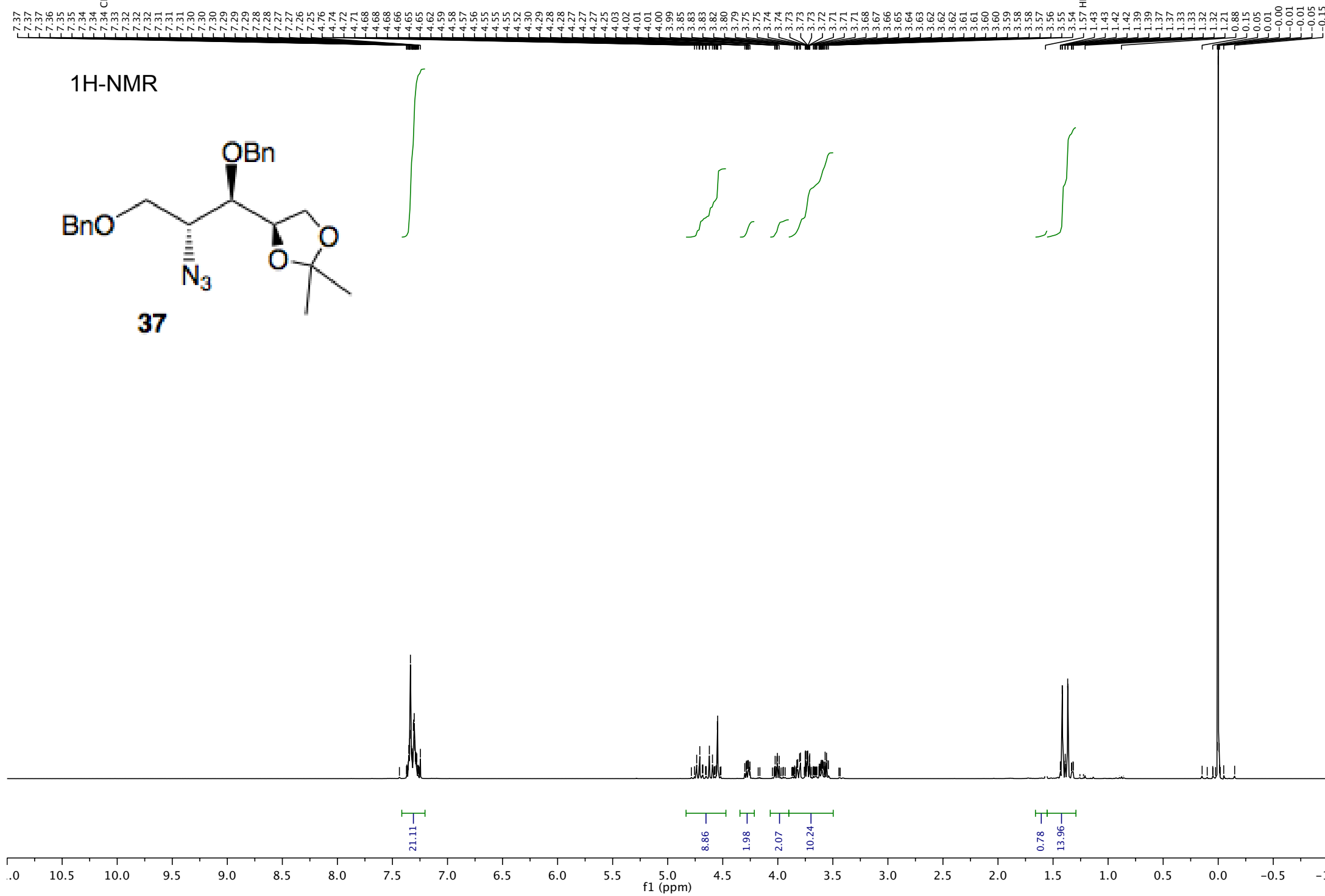
¹³C-NMR



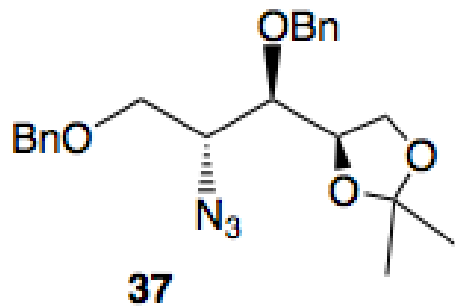
23

- 73.69
- 68.25
- 63.90
- 59.68
- 56.84
- 49.16
- 40.10
- 39.89 DMSO
- 39.68 DMSO
- 39.57 DMSO
- 39.26 DMSO
- 39.05 DMSO
- 28.87
- 28.75
- 28.59
- 26.55
- 26.45
- 22.05
- 13.90





13C-NMR



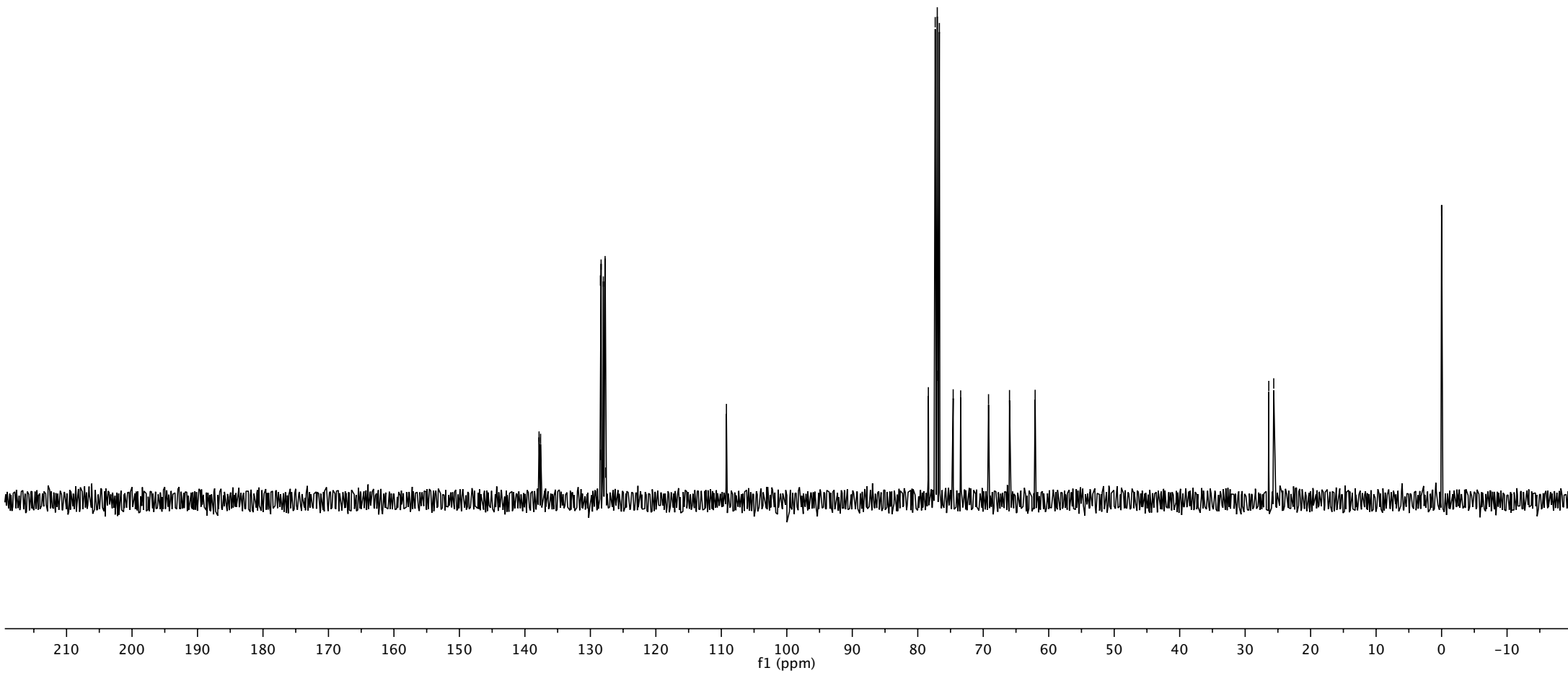
137.84
137.60
128.47
128.43
128.36
128.12
128.02
127.84
127.81
127.75
127.65

109.22

78.38
77.34 CDC13
77.02 CDC13
76.95 CDC13
76.70 CDC13
74.59
73.45
69.19
65.99
62.07

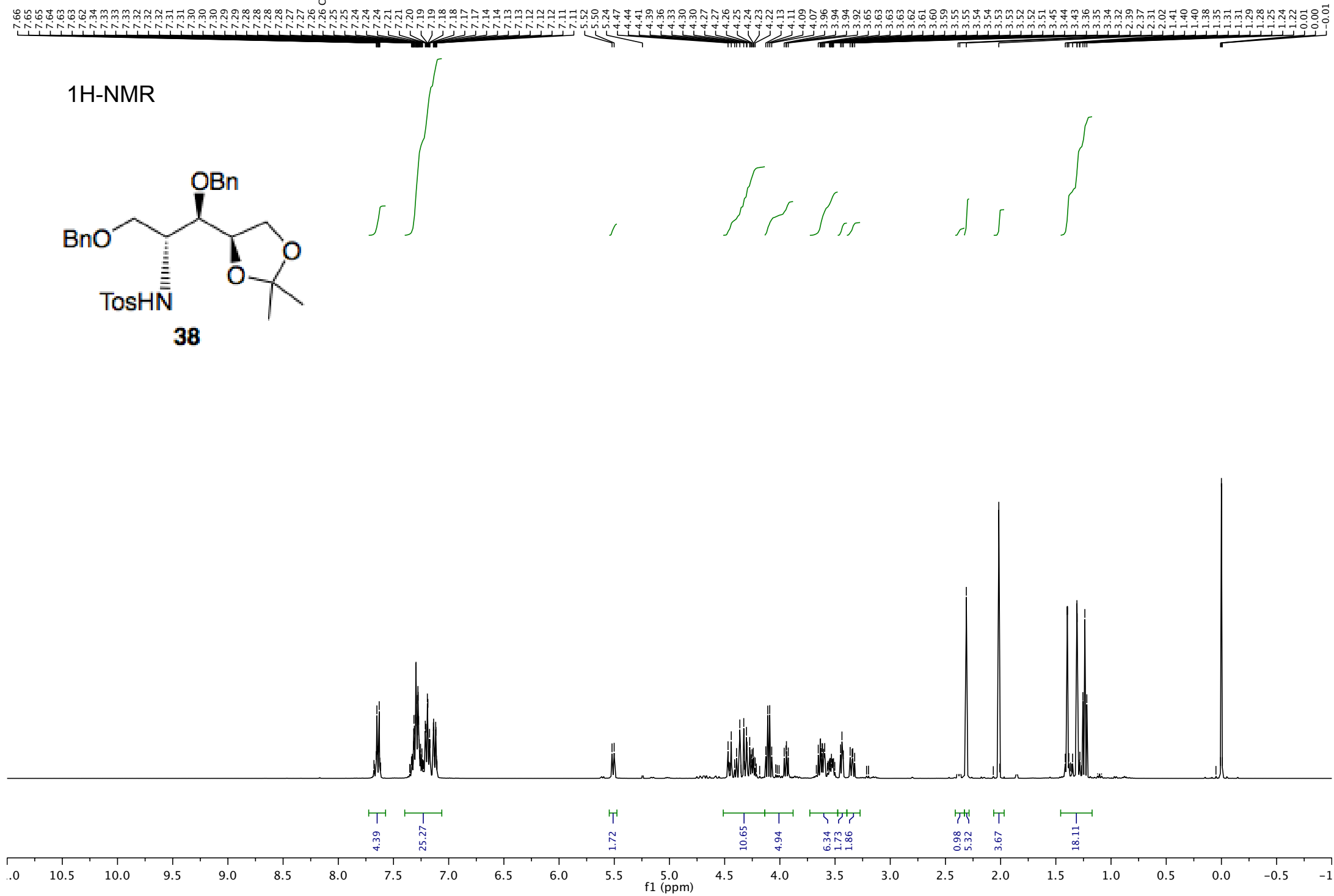
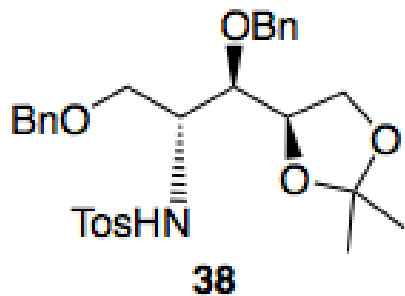
26.39
25.63

0.00

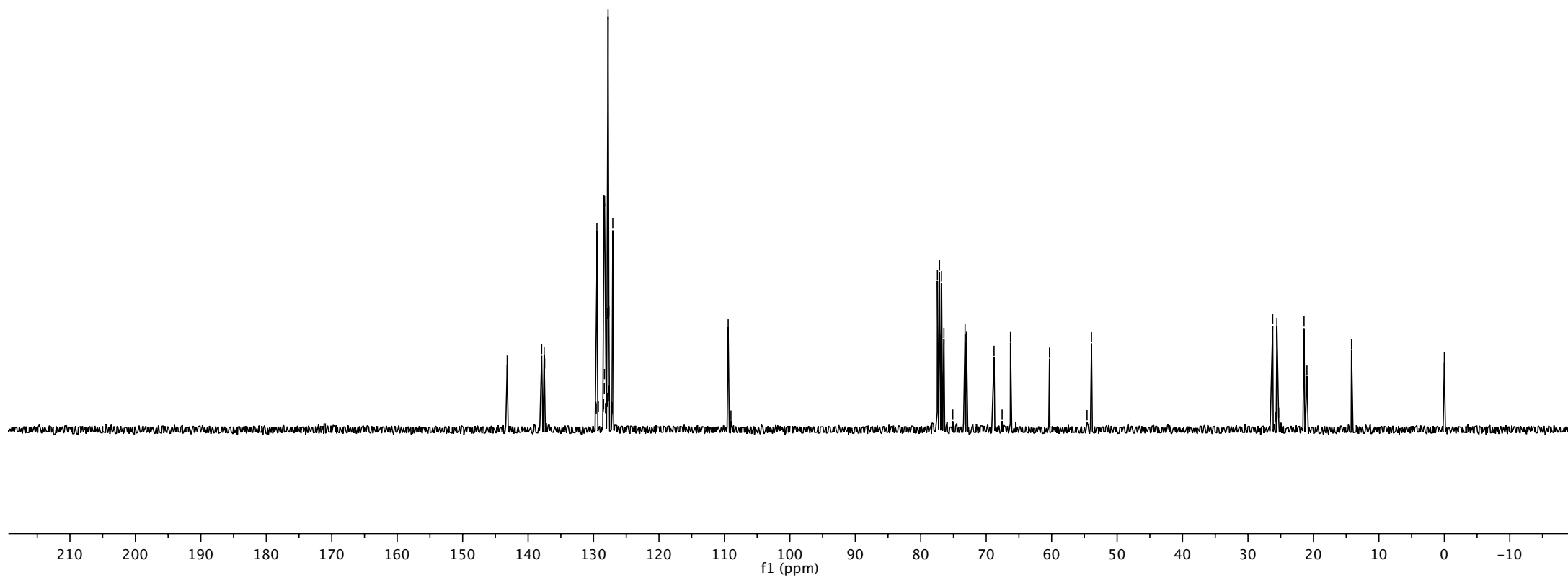
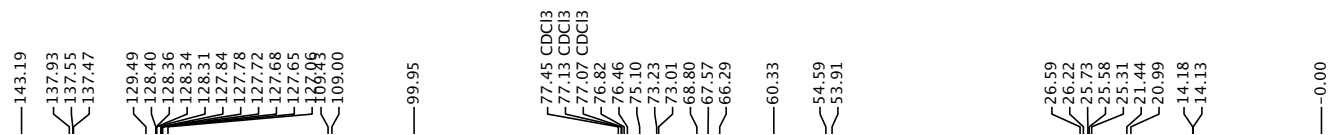
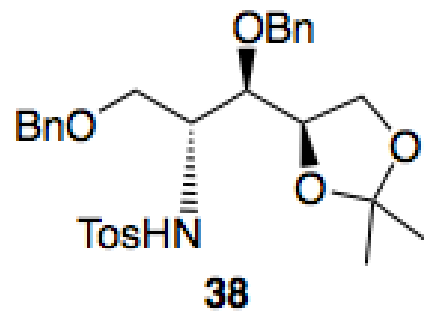


Aug29-2007
meso-tosyl

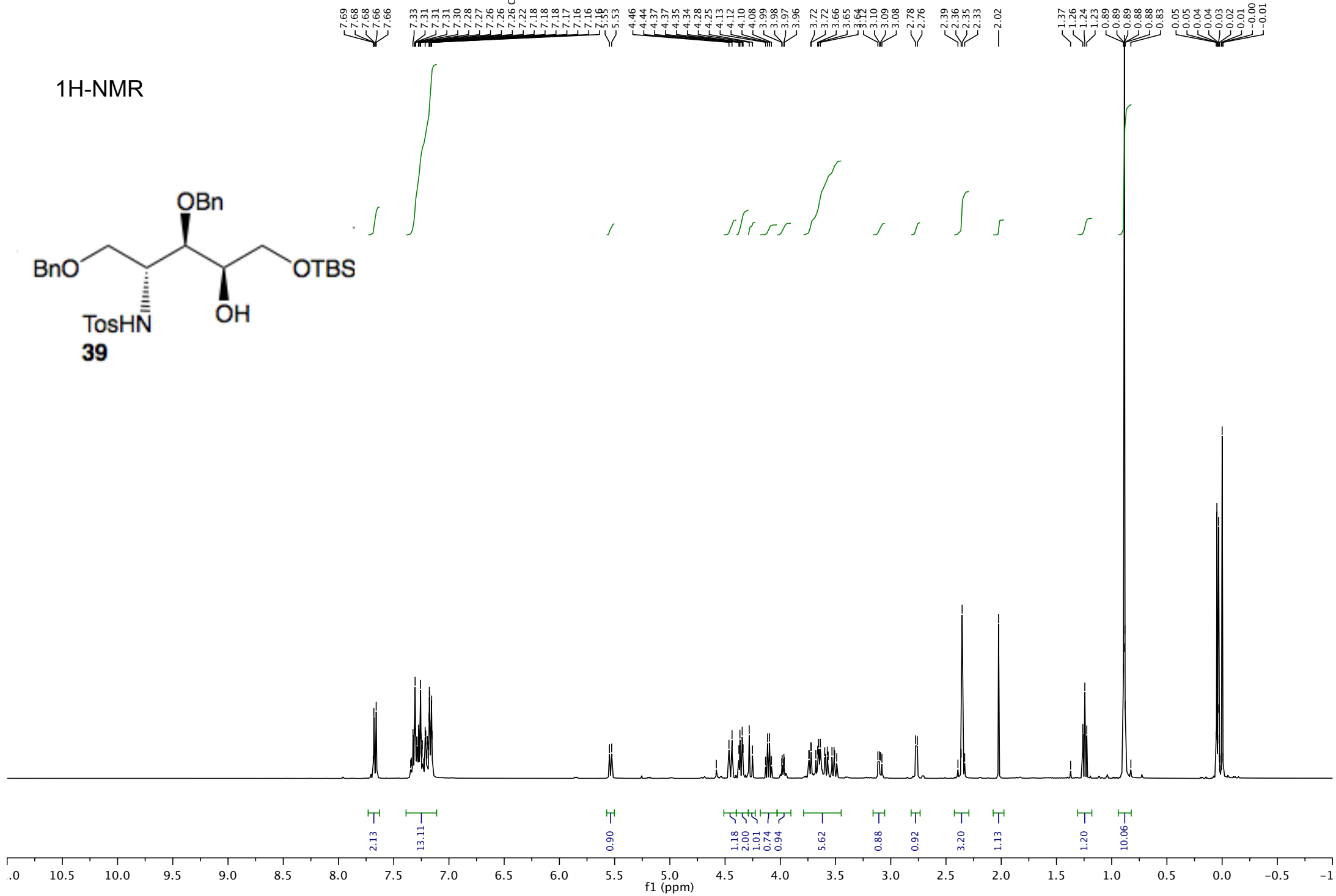
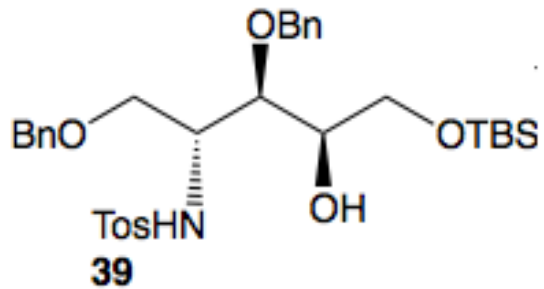
1H-NMR

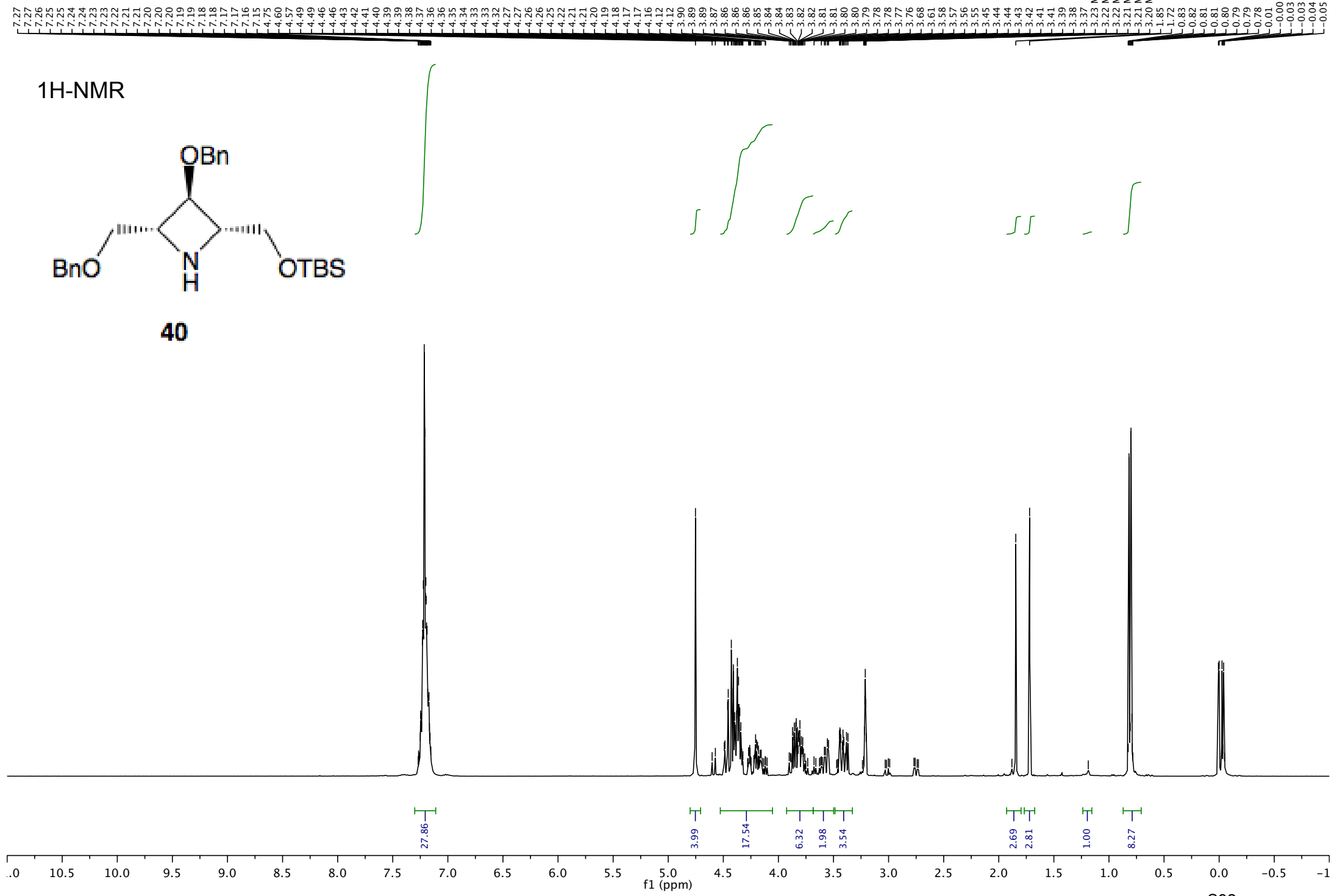


13C-NMR

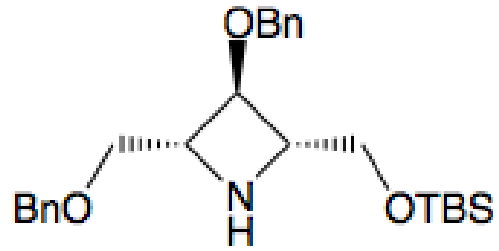


1H-NMR





13C-NMR



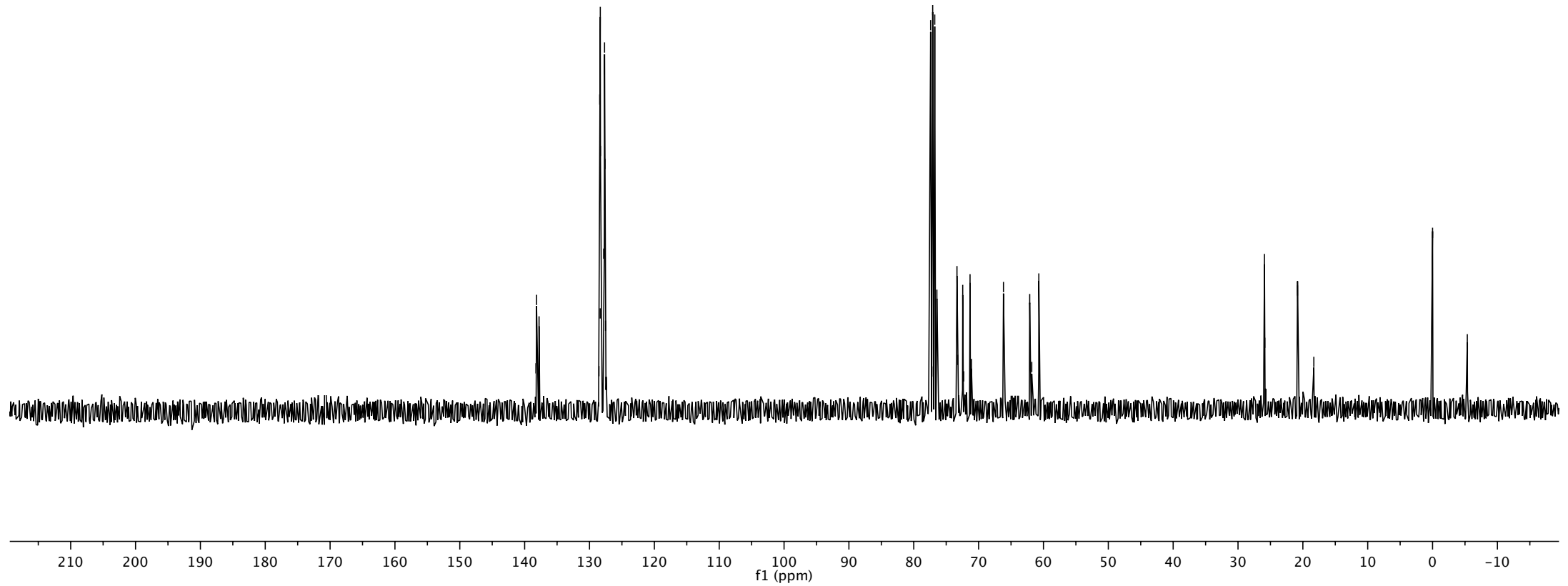
40

138.26
138.16
137.78
128.54
128.42
128.35
128.34
128.31
127.82
127.71
127.68
127.65
127.60
127.53

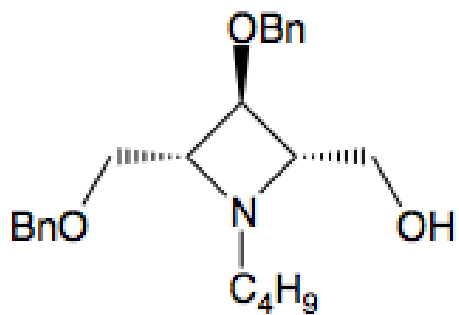
77.40 CDCl3
77.08 CDCl3
76.98 CDCl3
76.76 CDCl3
76.43
73.34
72.42
72.45
72.30
71.32
71.10
66.17
64.93
63.93
62.12
61.80
60.71

25.92
25.87
20.82
18.31

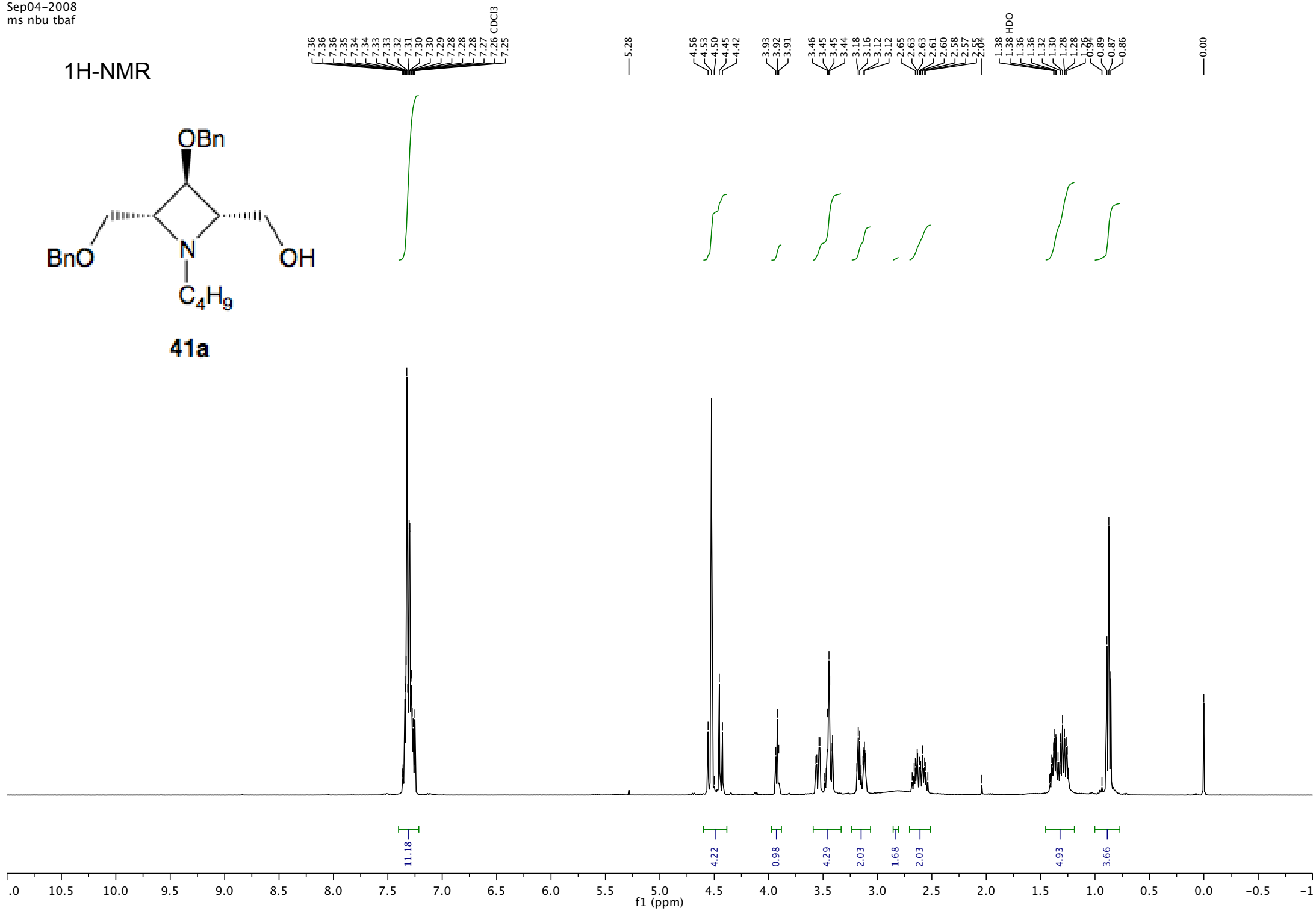
-0.00
-5.36
-5.38



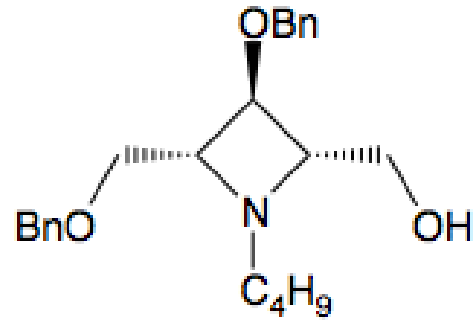
1H-NMR



41a



13C-NMR

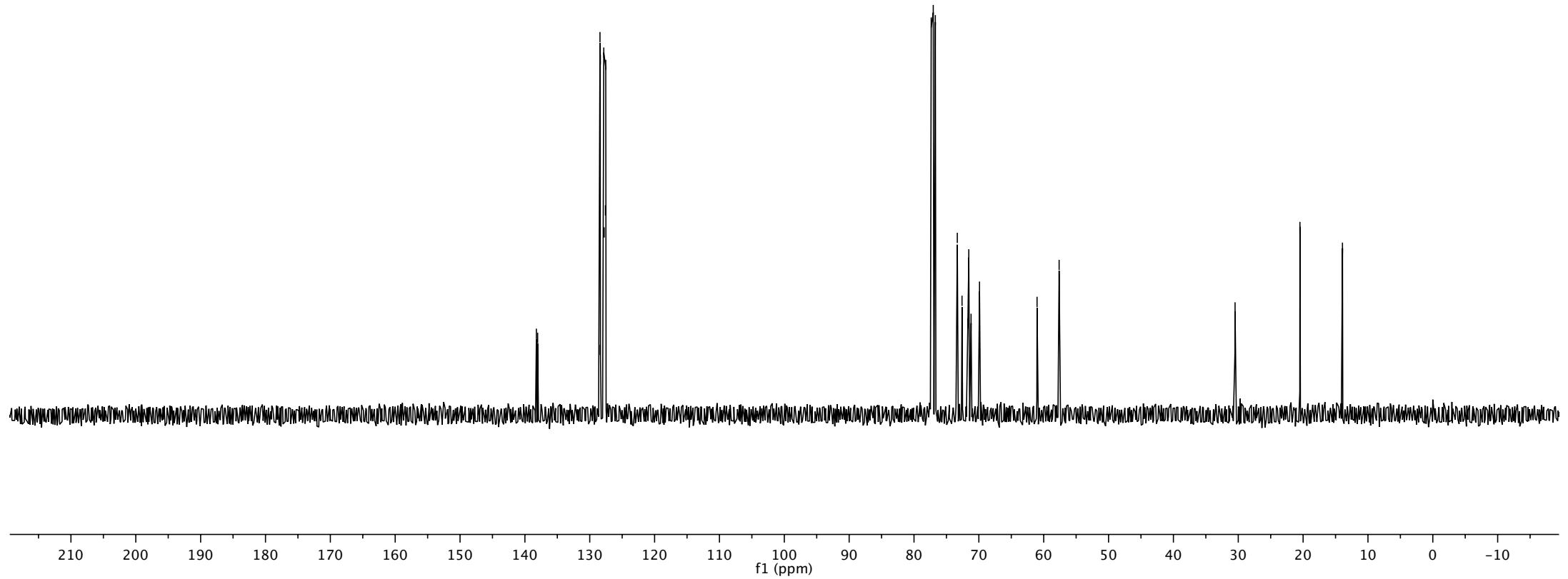


41a

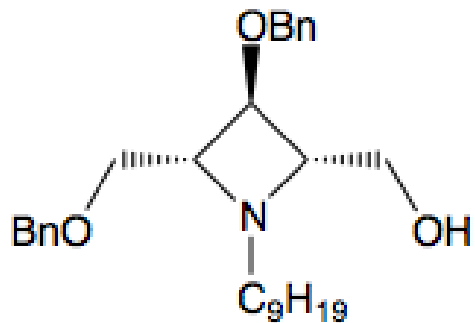
138.23
138.03
128.42
128.38
128.36
127.83
127.78
127.58
127.53

77.35 CDC13
77.03 CDC13
76.71 CDC13
73.32
72.60
71.71
71.55
71.20
69.90
61.03
57.62

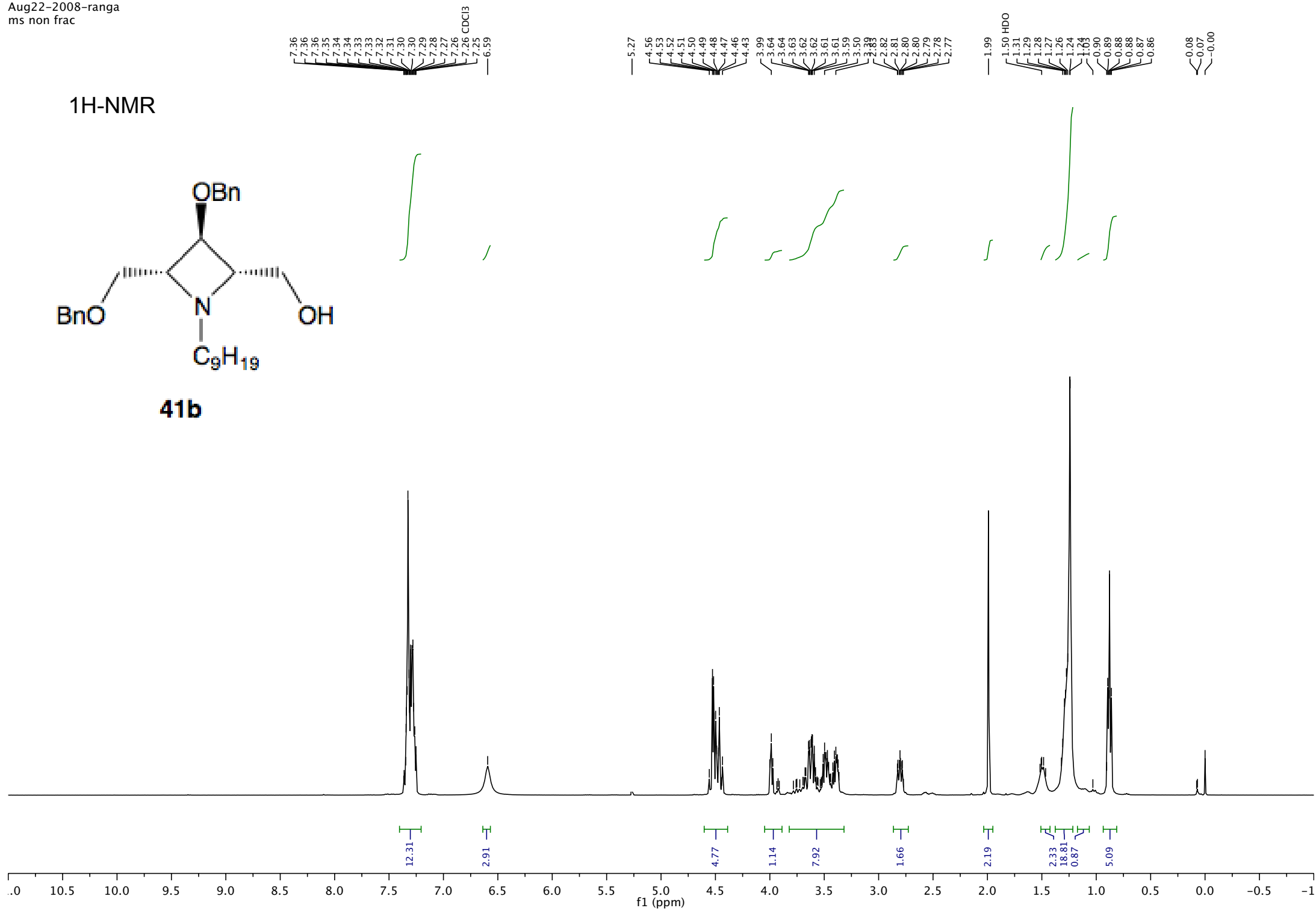
30.49
20.49
13.93



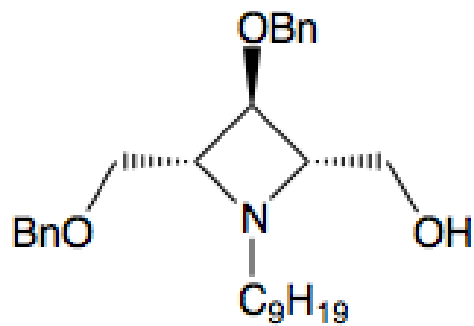
1H-NMR



41b



13C-NMR



41b

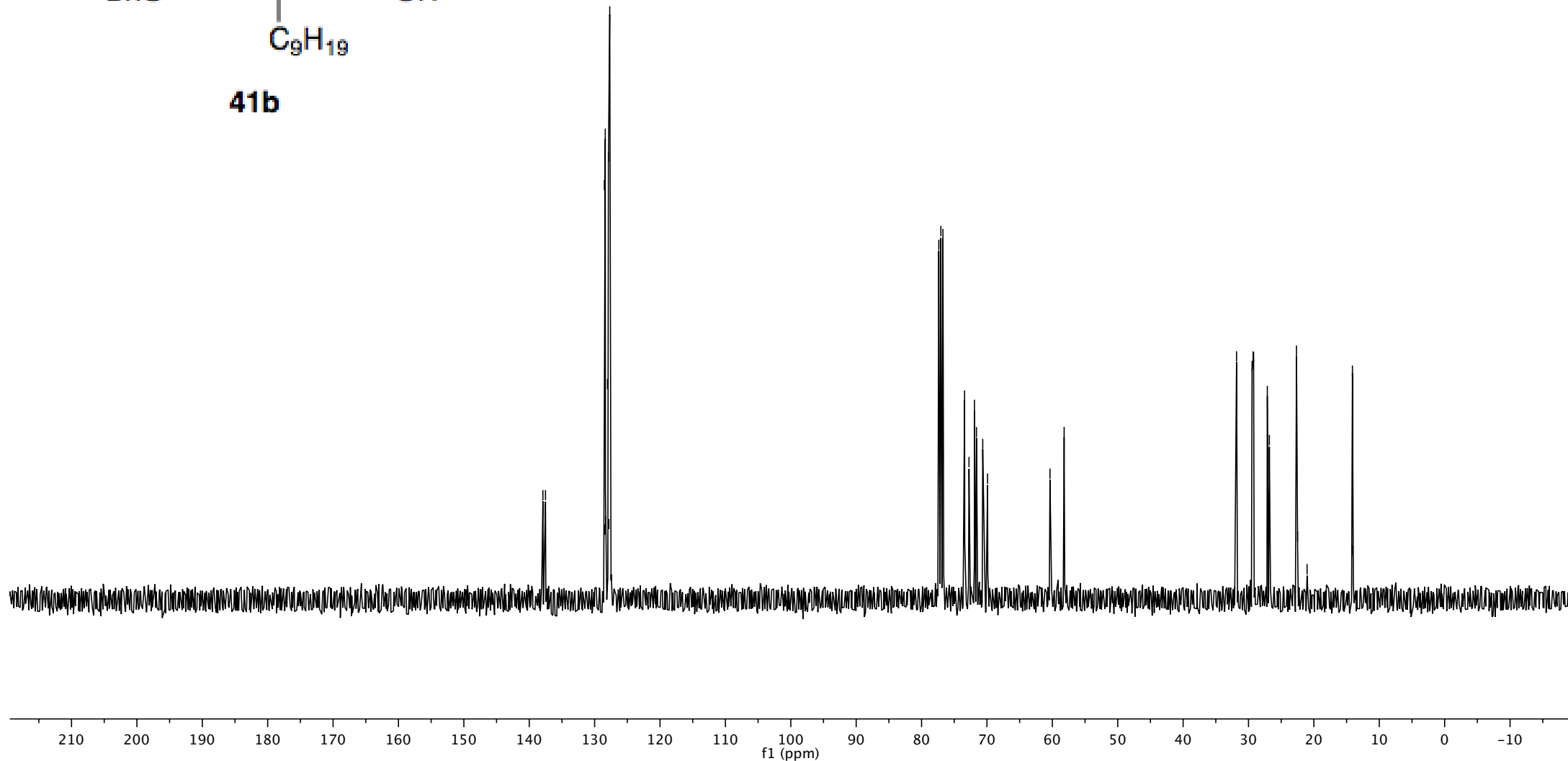
137.91
137.52
128.51
128.43
128.38
128.32
128.02
127.89
127.82
127.69

96.43

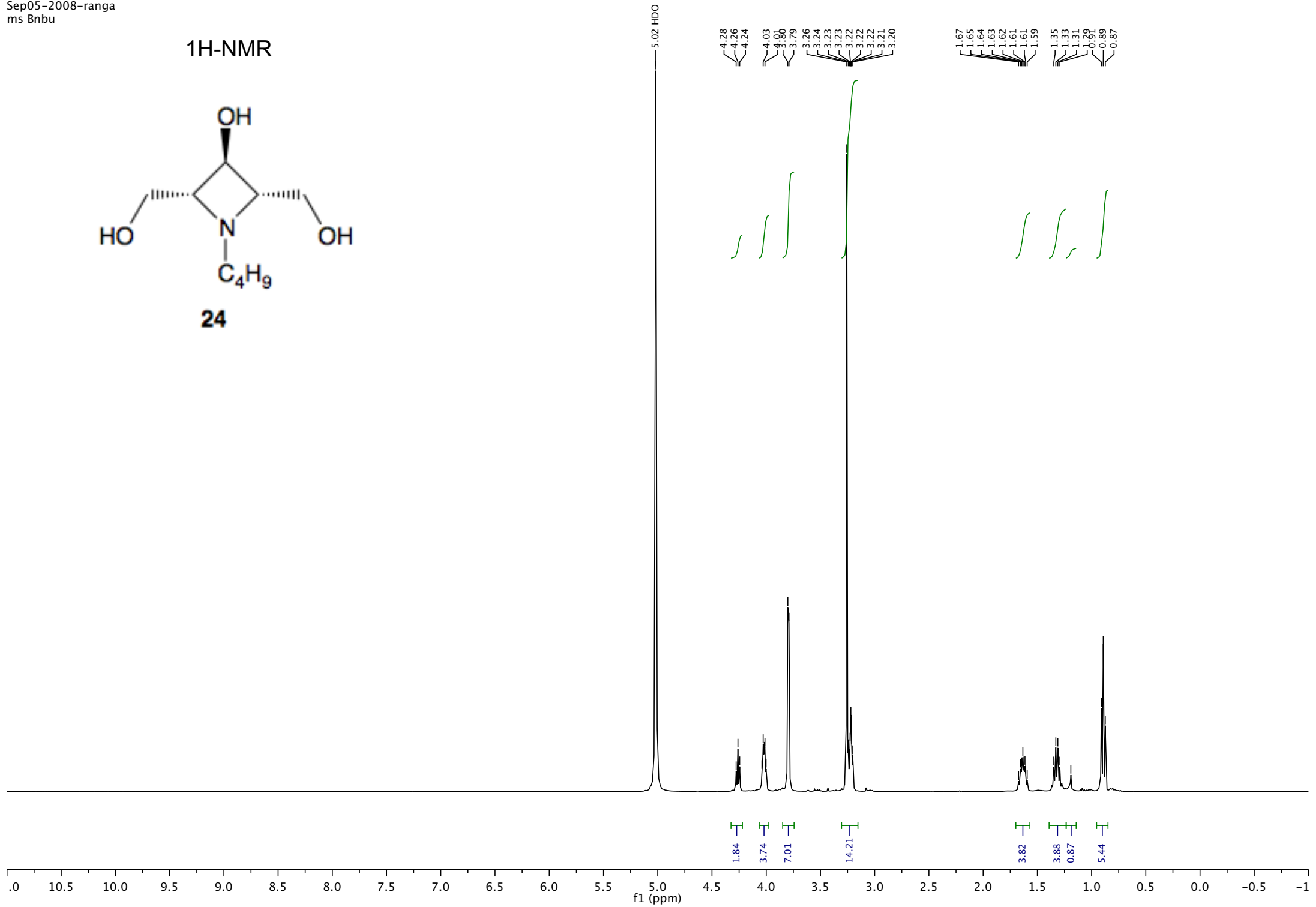
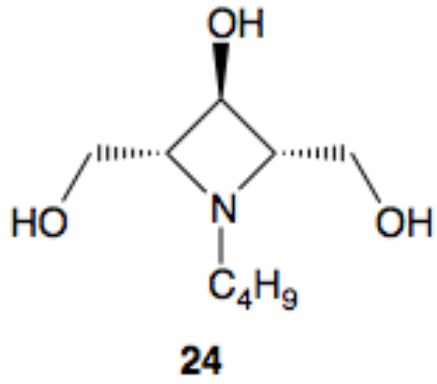
77.37 CDCl₃
77.05 CDCl₃
76.73 CDCl₃
73.42
72.74
71.91
71.60
70.64
69.90

60.36
58.19

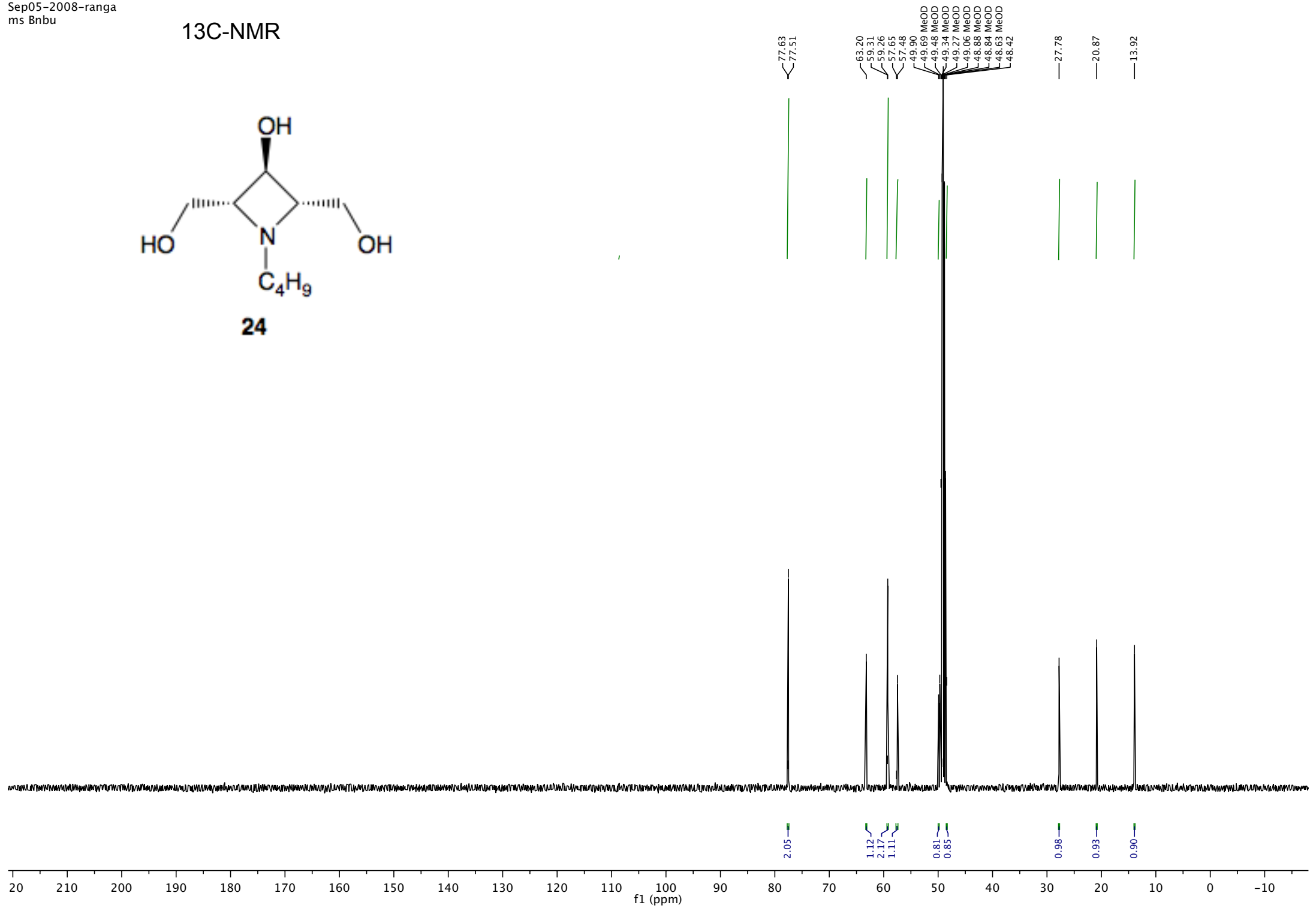
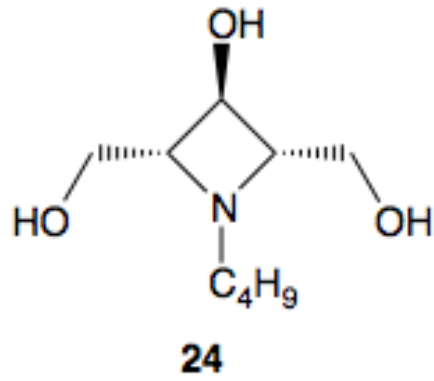
31.83
29.46
29.30
29.22
27.12
26.81
22.66
22.49
21.04
14.19
14.11



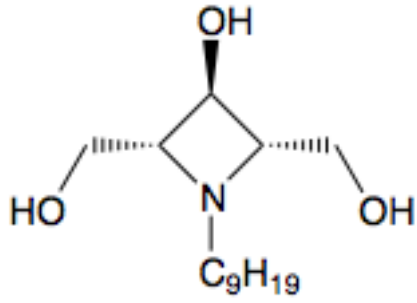
1H-NMR



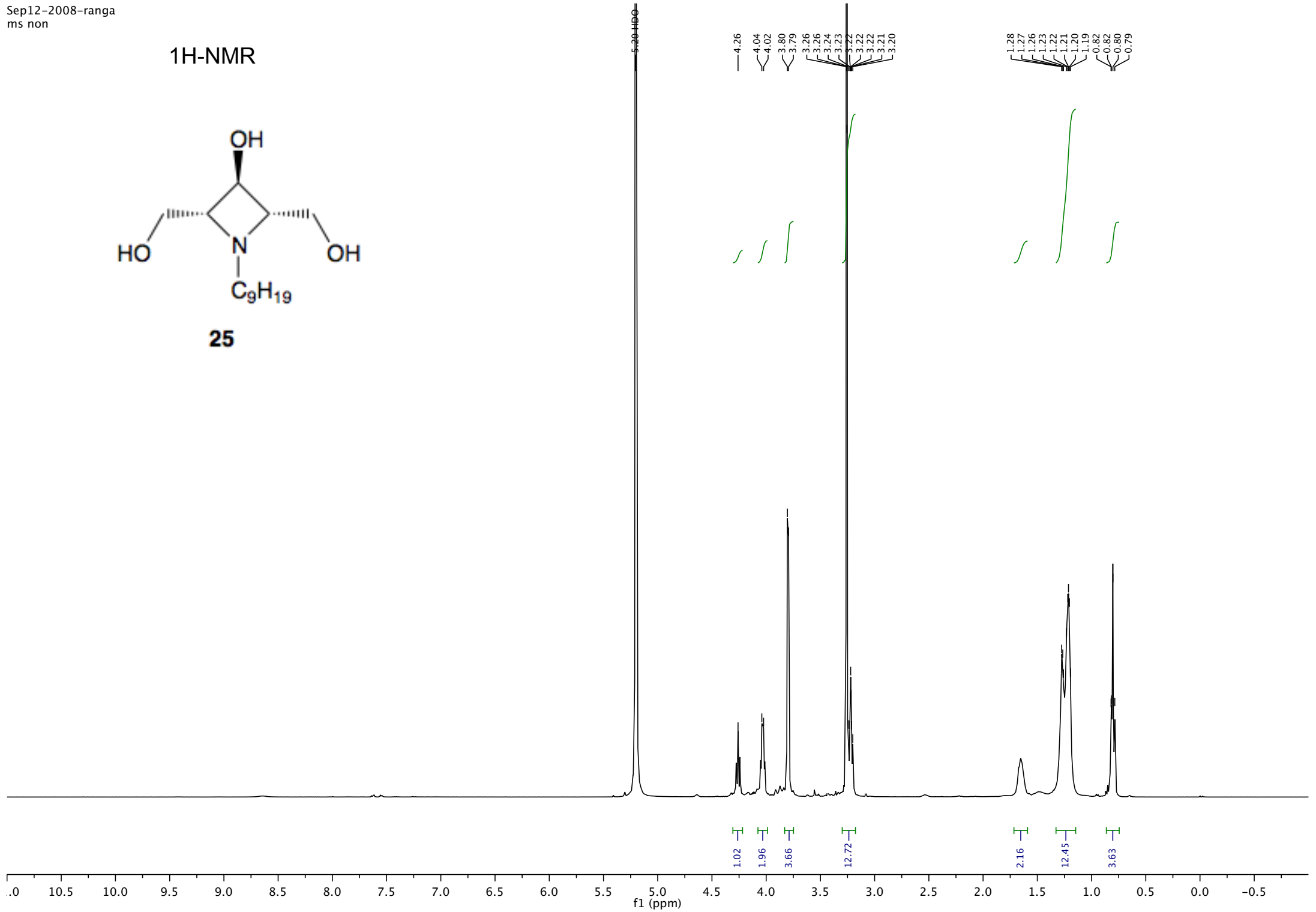
13C-NMR



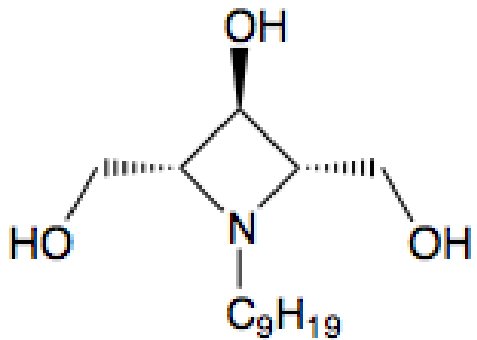
1H-NMR



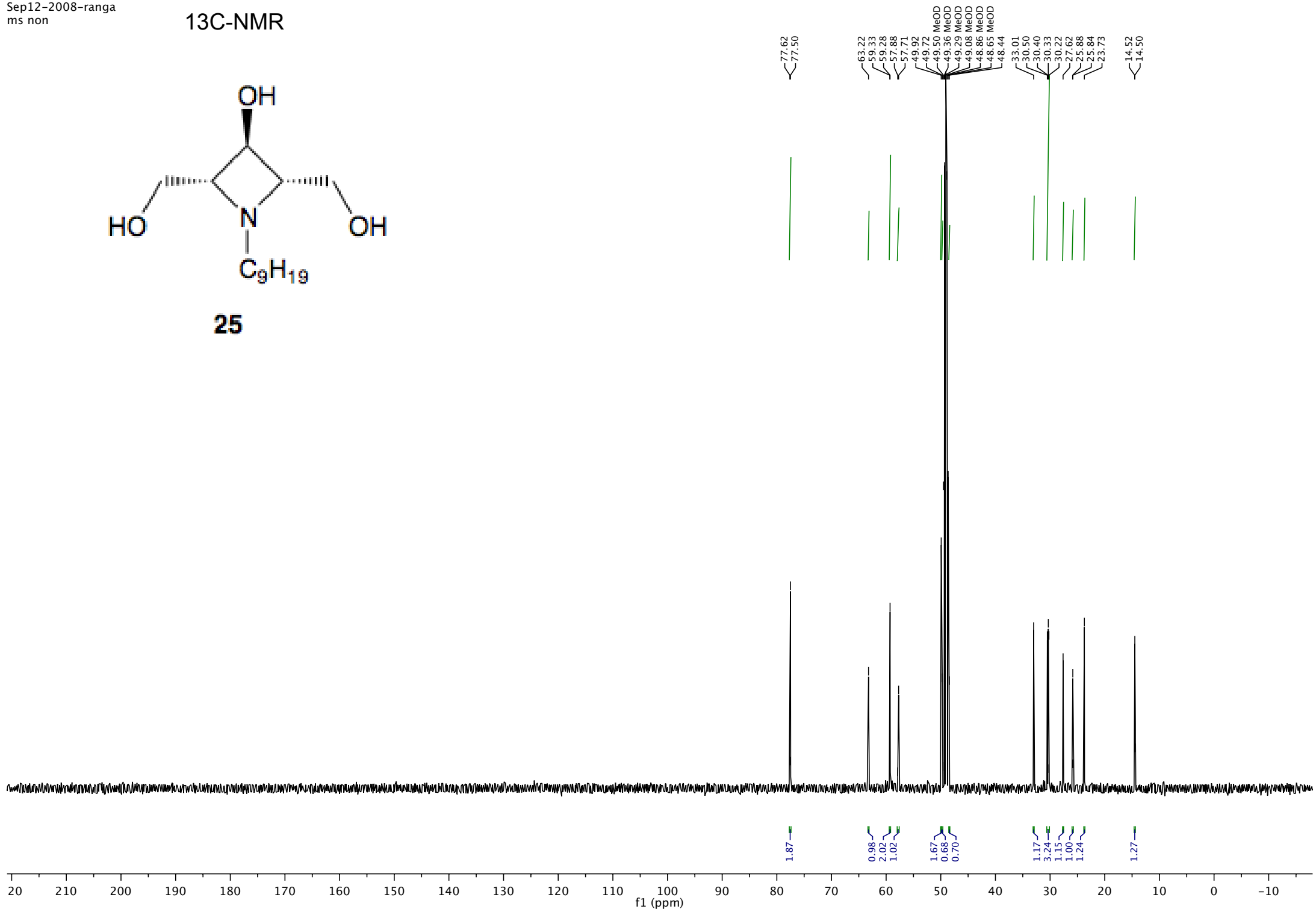
25



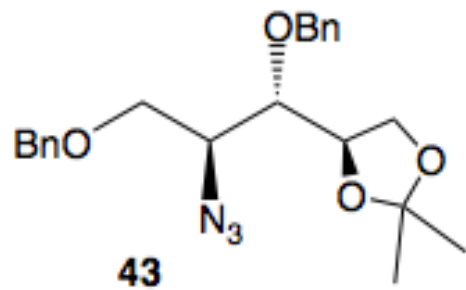
13C-NMR



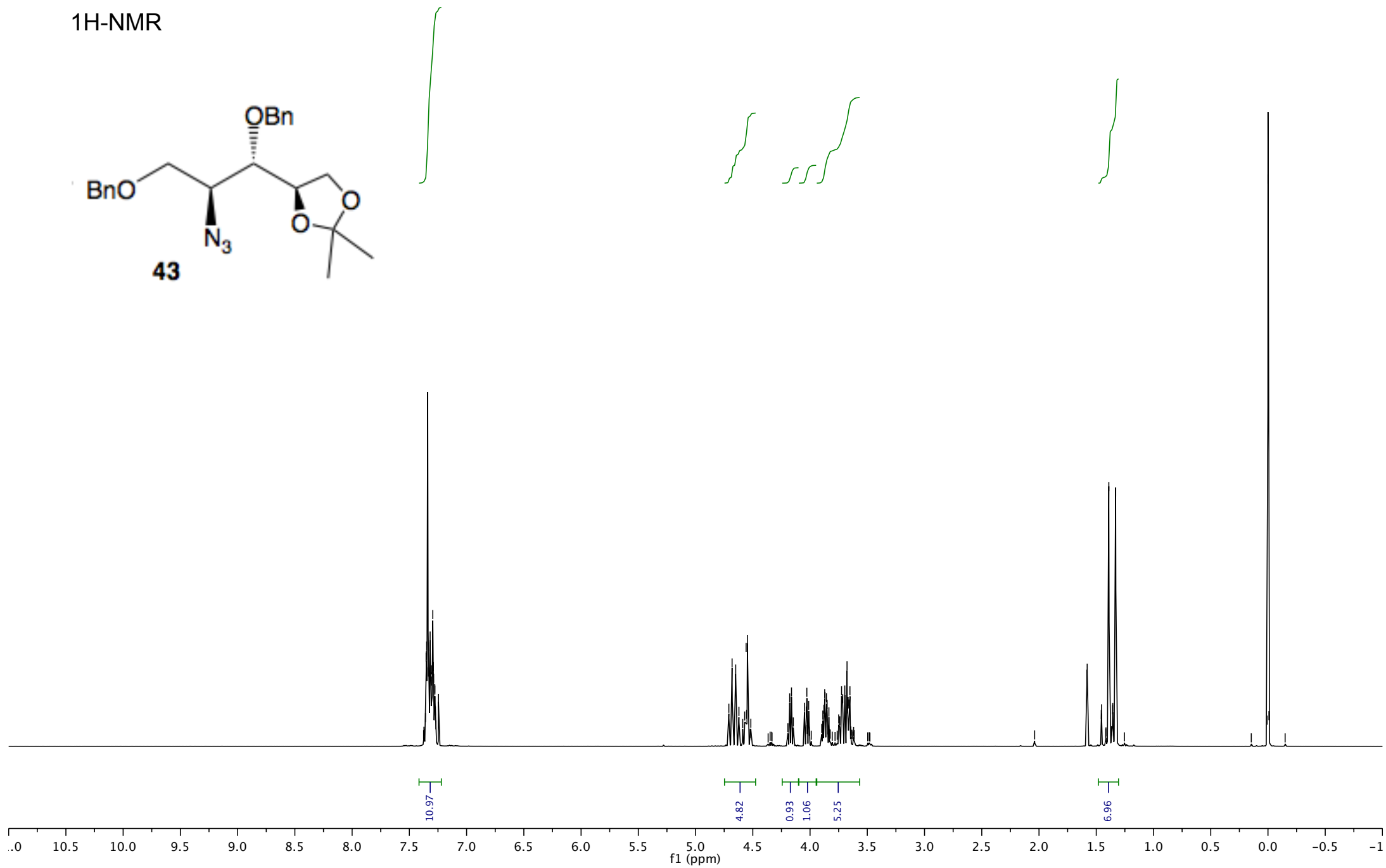
25



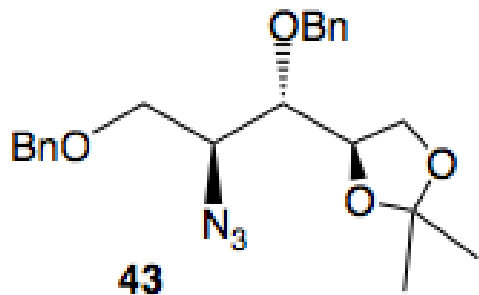
1H-NMR



7.38
7.37
7.36
7.35
7.35
7.35
7.34 CDCl3
7.33
7.32
7.31
7.30
7.29
7.29
7.28
7.28
7.27
7.25
4.71
4.68
4.65
4.62
4.59
4.57
4.56
4.55
4.55
4.52
4.19
4.18
4.16
4.15
4.05
4.03
4.02
4.01
3.90
3.89
3.88
3.87
3.86
3.86
3.85
3.85
3.84
3.75
3.74
3.72
3.71
3.70
3.68
3.67
3.67
3.66
3.65
2.64 H2O
1.58 H2O
1.46
1.45
1.42
1.39
1.39
1.37
1.37
1.36
1.36
1.33
1.25
0.15
0.01
-0.01
-0.15



13C-NMR



137.72
137.65

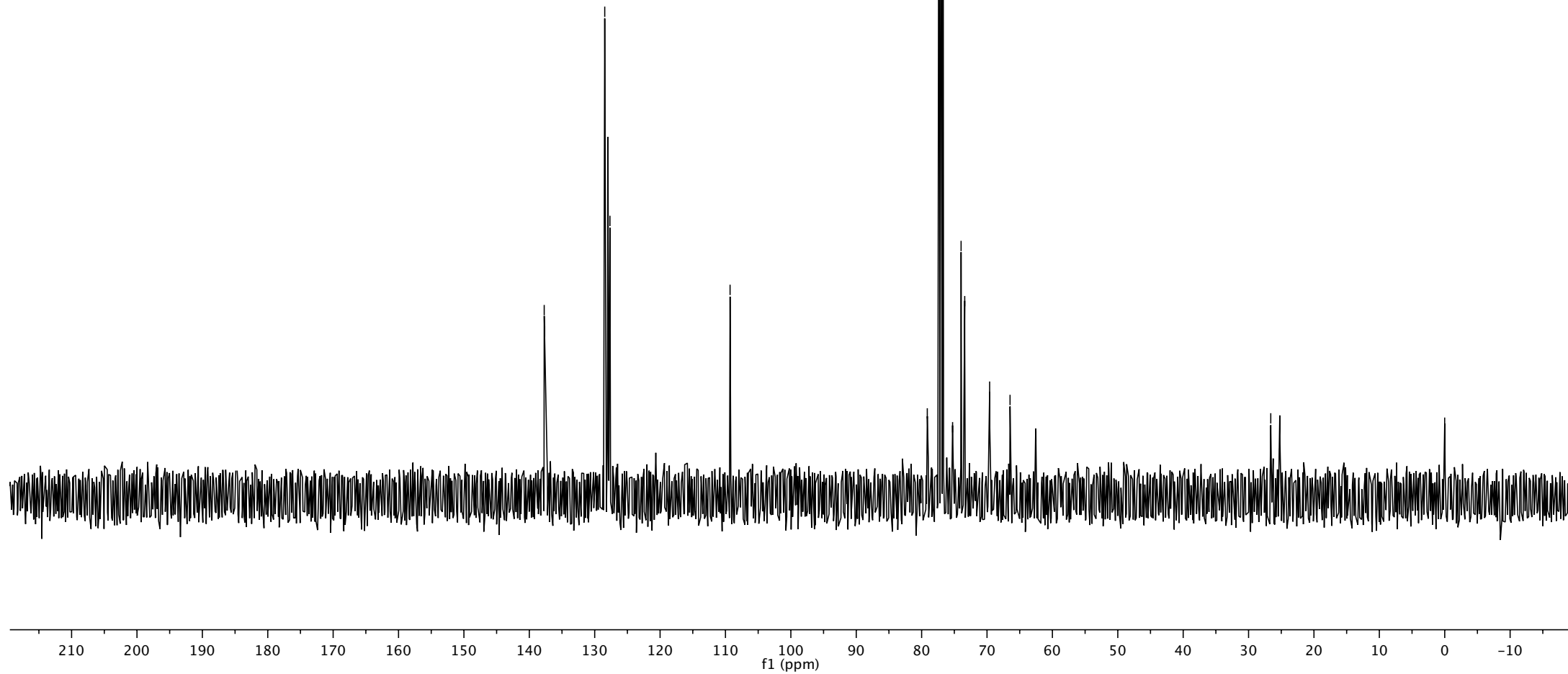
128.46
127.97
127.67

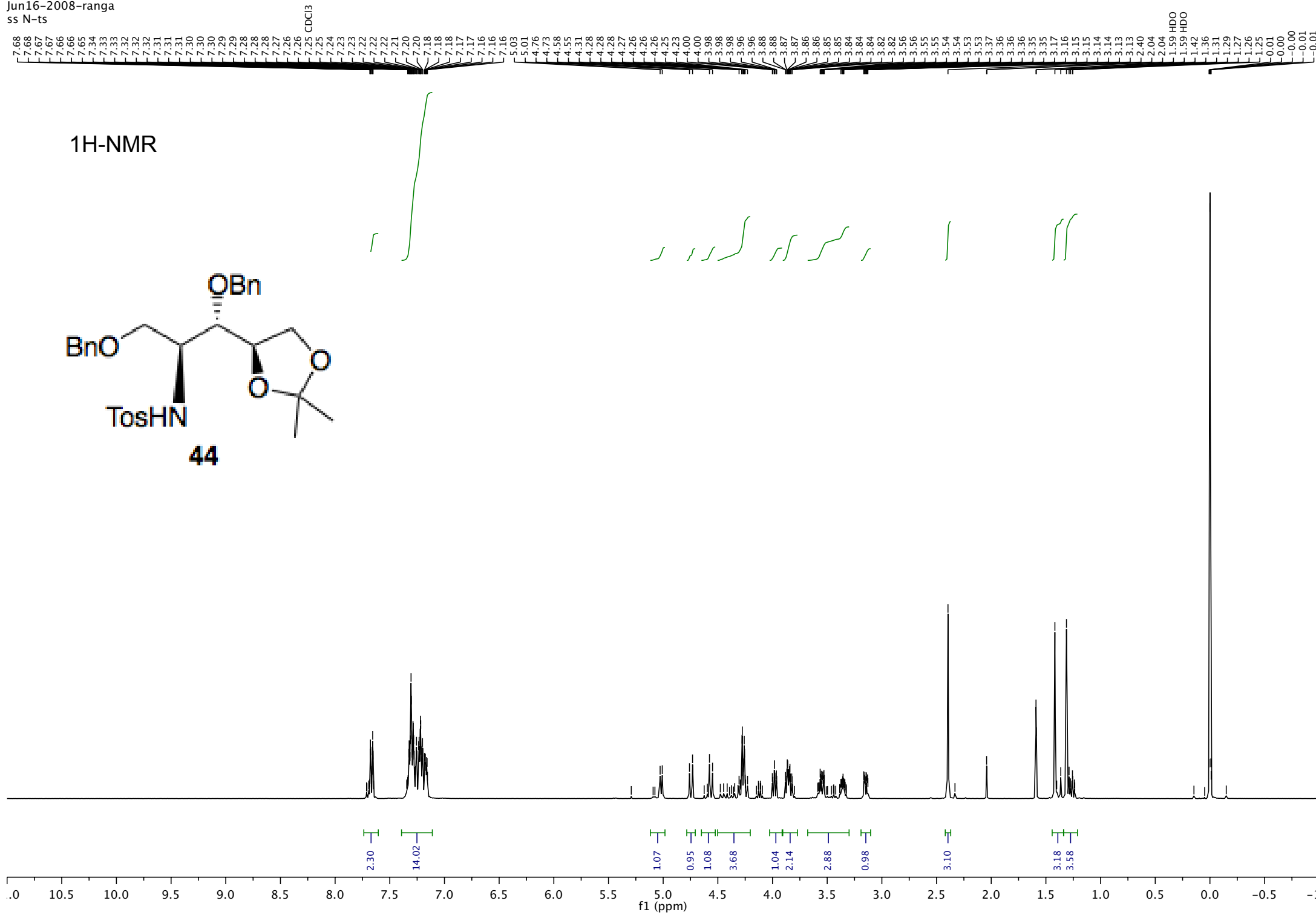
109.29

79.13
77.35 CDCl3
77.04 CDCl3
76.72 CDCl3
75.27
73.97
73.41
69.60
66.48
62.52

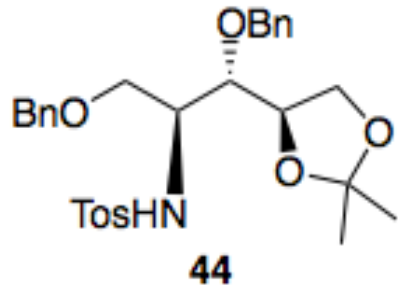
26.61
25.25

0.01





13C-NMR



143.50
138.06
137.46
137.15
129.63
128.42
128.38
128.36
127.90
127.87
127.78
127.72
127.18

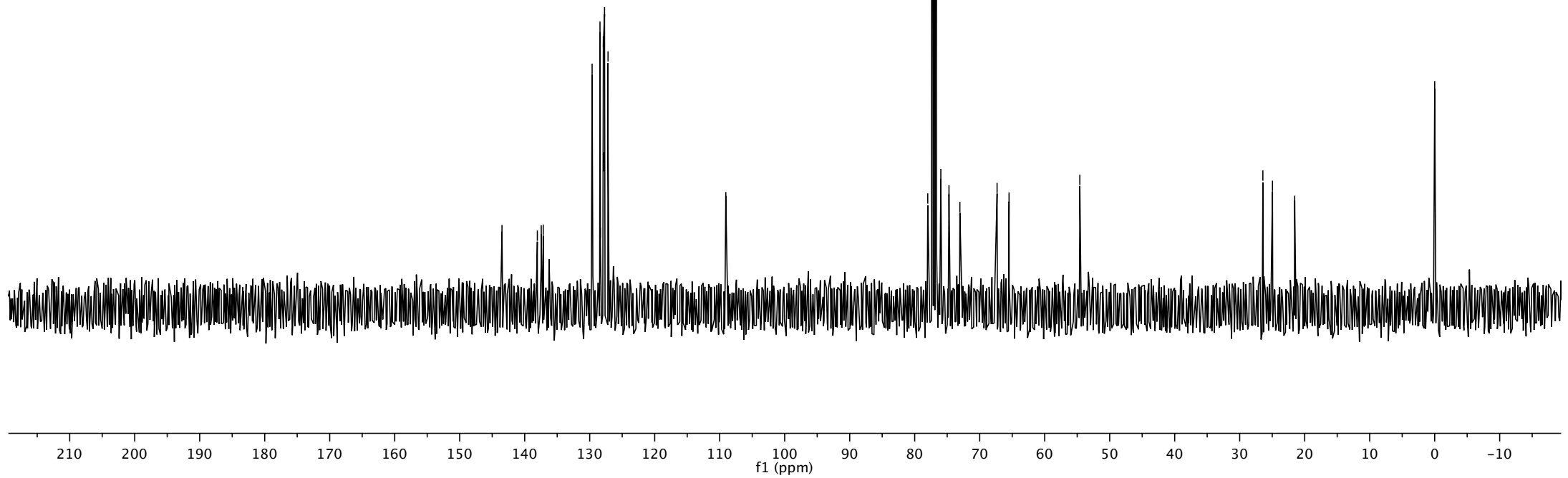
109.06

77.99 CDCI3
77.34 CDCI3
77.02 CDCI3
76.70 CDCI3
75.98
74.73
73.05
67.33
65.48

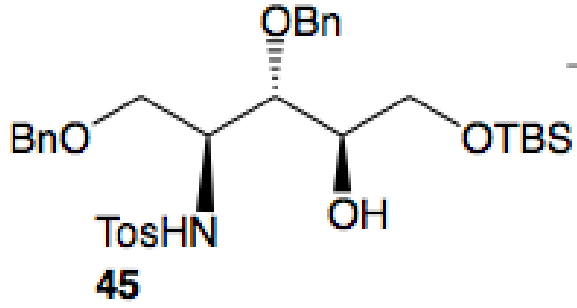
54.60

26.43
24.96
21.54

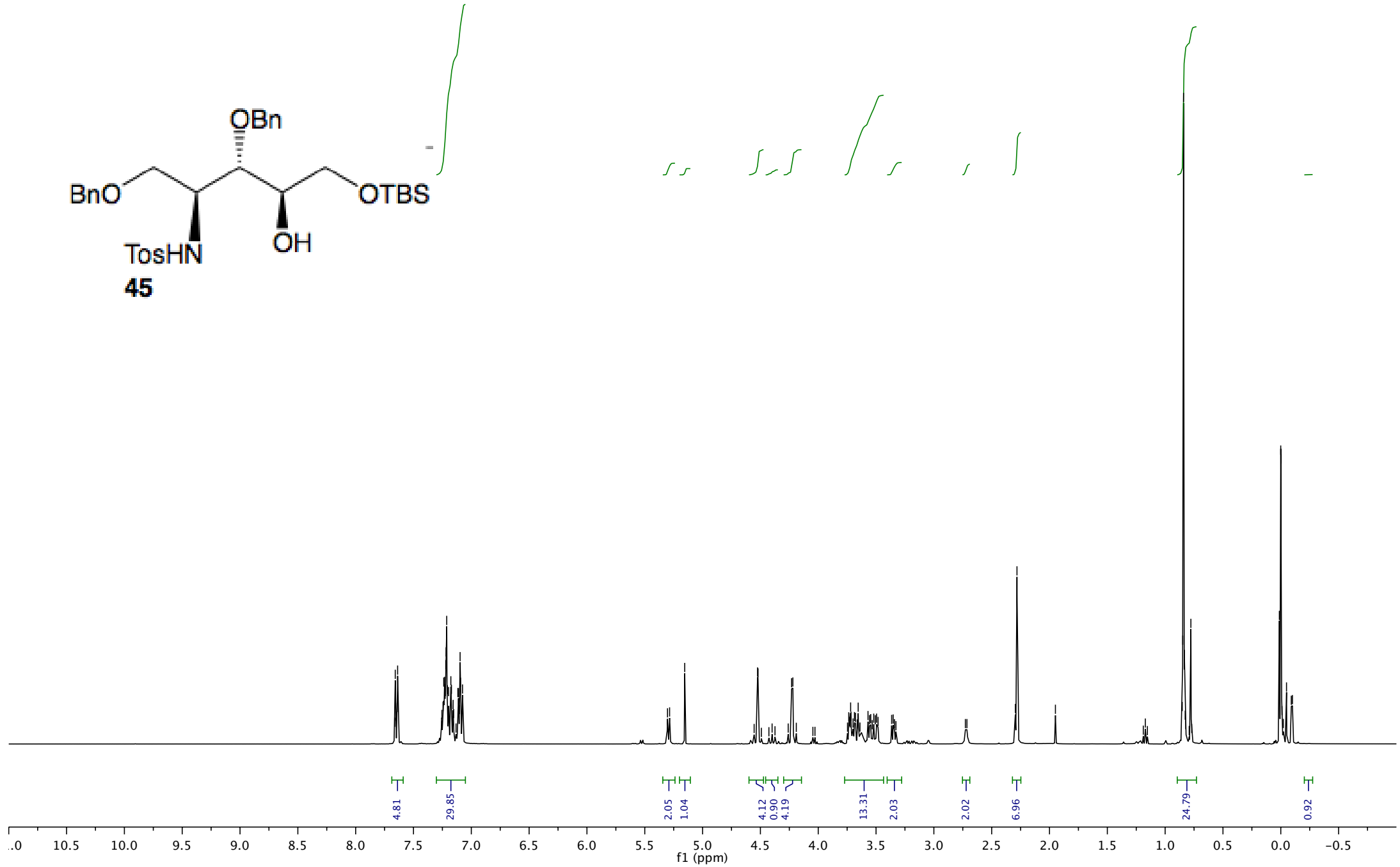
-0.00



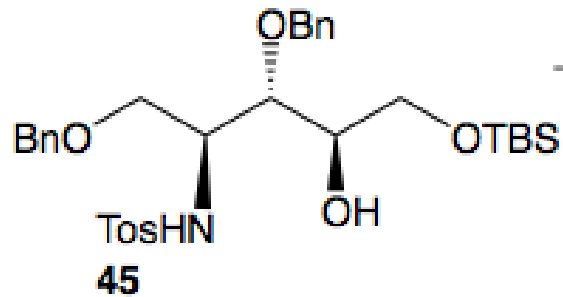
1H-NMR



7.66
7.65
7.64
7.64
7.24
7.23
7.22
7.22
7.22
7.21
7.21
7.20
7.18
7.17
7.16
7.15
7.12
7.11
7.10
7.10
7.09
5.98
5.29
5.15
4.55
4.52
4.52
4.43
4.40
4.37
4.26
4.23
4.22
4.19
4.05
4.03
3.74
3.72
3.72
3.69
3.68
3.65
3.57
3.55
3.52
3.52
3.59
2.71
2.30
2.28
1.95
1.19
1.17
1.15
0.85
0.85
0.84
0.83
0.78
0.78
0.00
-0.01
-0.02
-0.05
-0.09
-0.10



13C-NMR

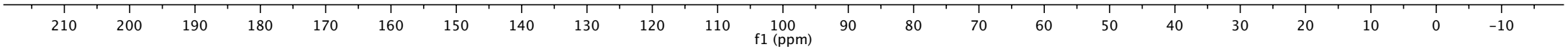


143.05
143.02
138.04
137.59
136.54
129.50
129.44
128.40
128.37
128.33
128.27
128.16
127.90
127.88
127.84
127.70
127.63
127.57
127.55
127.15
127.03

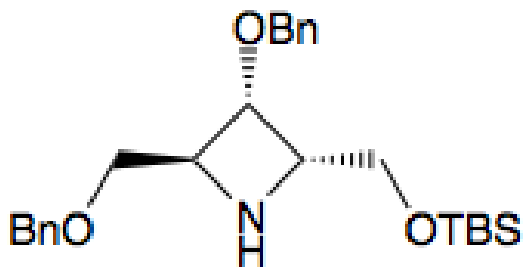
80.13
78.56
77.46
77.14 CDCl3
76.82
73.74
73.21
73.17
72.90
71.53
70.22
69.61
68.29
63.67
63.60
60.31

25.86
25.78
25.66
21.45
21.43
18.19
18.08
17.93
14.15

-0.00
-3.58
-5.40
-5.43
-5.53
-5.55



1H-NMR



46

7.30
7.29
7.28
7.27
7.27 CDC13
7.26
7.25
7.24
7.22
7.22
7.21
7.20
7.19

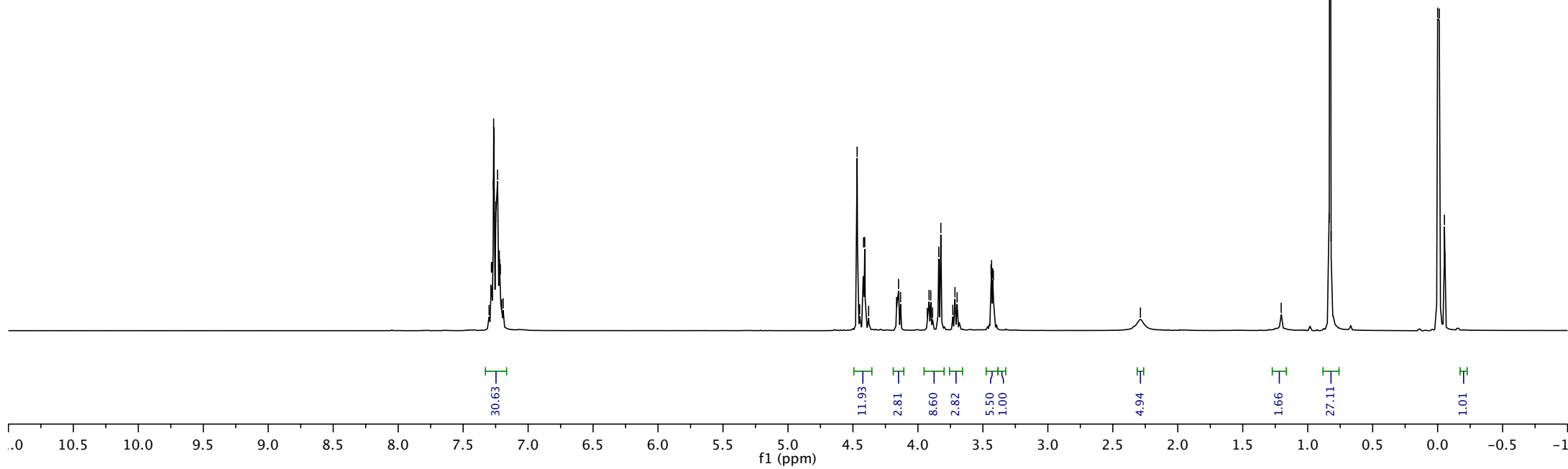
4.47
4.45
4.42
4.41
4.38
4.15
4.13
3.91
3.84
3.82
3.77
3.74
3.43
3.42
3.42

2.29

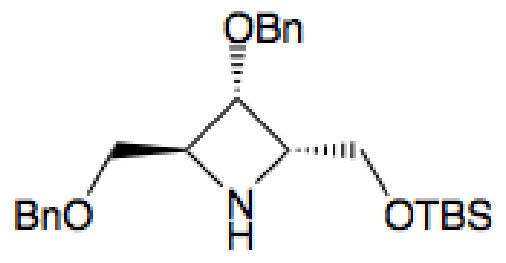
1.20

0.84
0.83
0.82

-0.00
-0.01
-0.05
-0.05



13C-NMR



46

138.17
138.09

128.35
128.33
127.68
127.64
127.61

77.37
77.05 CDCl3
76.73 CDCl3

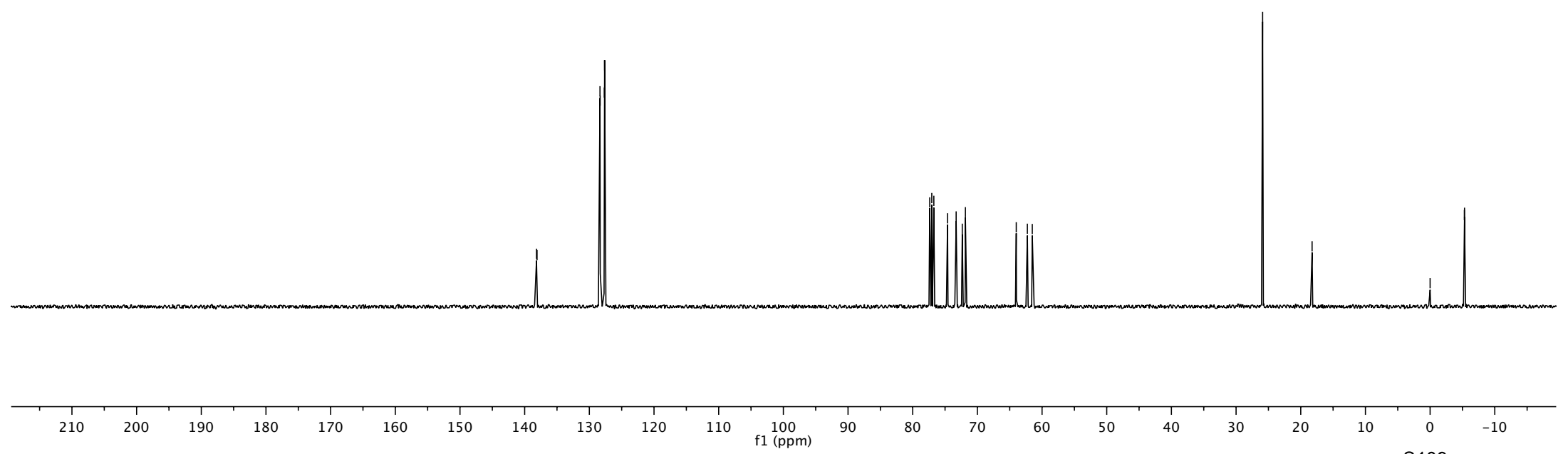
74.62
73.28
72.34
71.85

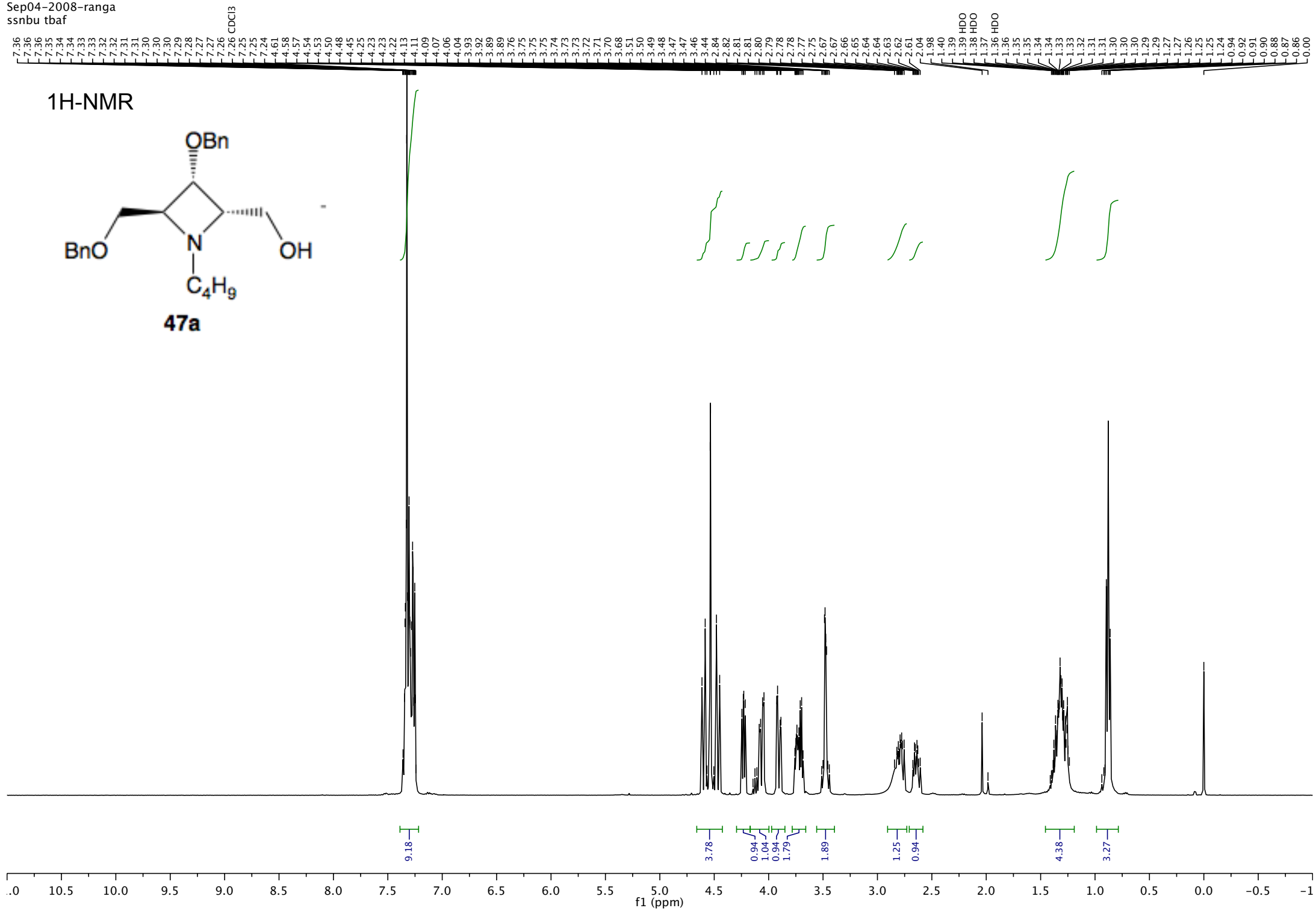
63.99
62.27
61.51

29.69
25.90

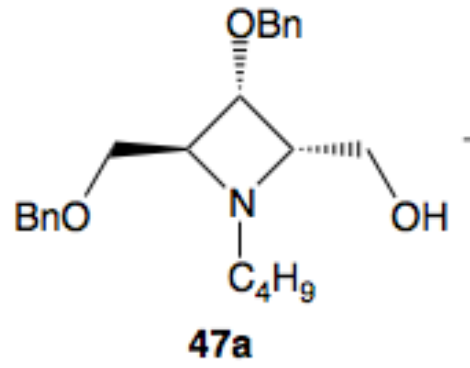
18.25

0.00
-5.30
-5.32





13C-NMR



138.20
137.63
128.48
128.36
127.87
127.69
127.68
127.63

77.36 CDCl3
77.04 CDCl3
76.72 CDCl3
74.86
73.36
71.91
71.50
71.09
64.92
60.37

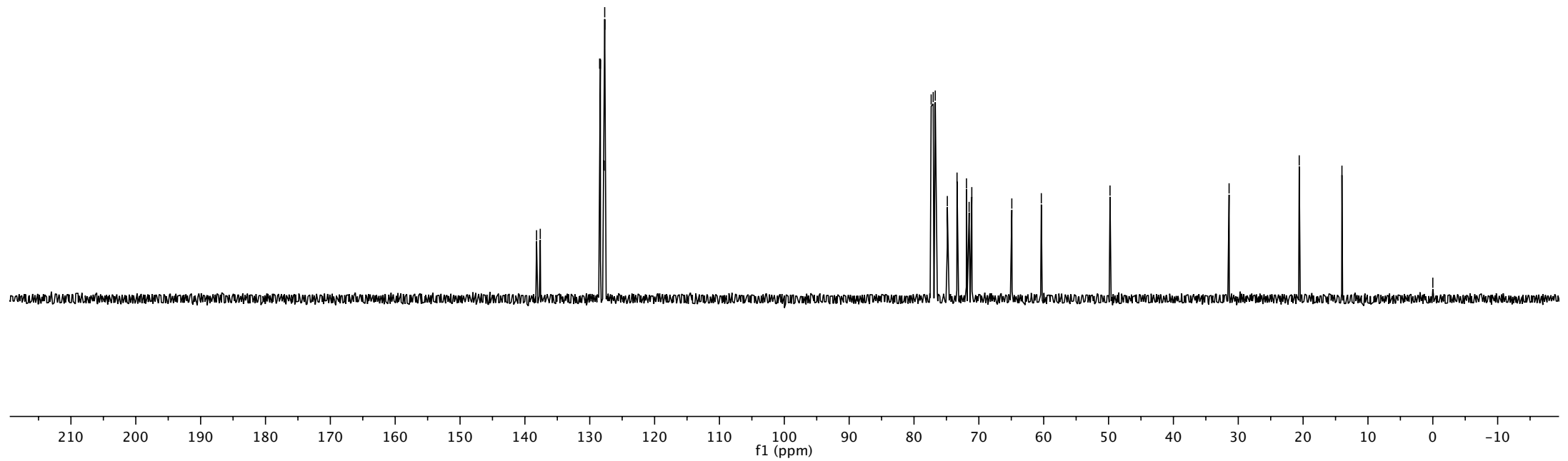
49.77

31.41

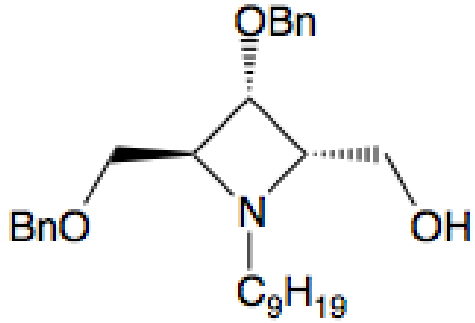
20.60

14.01

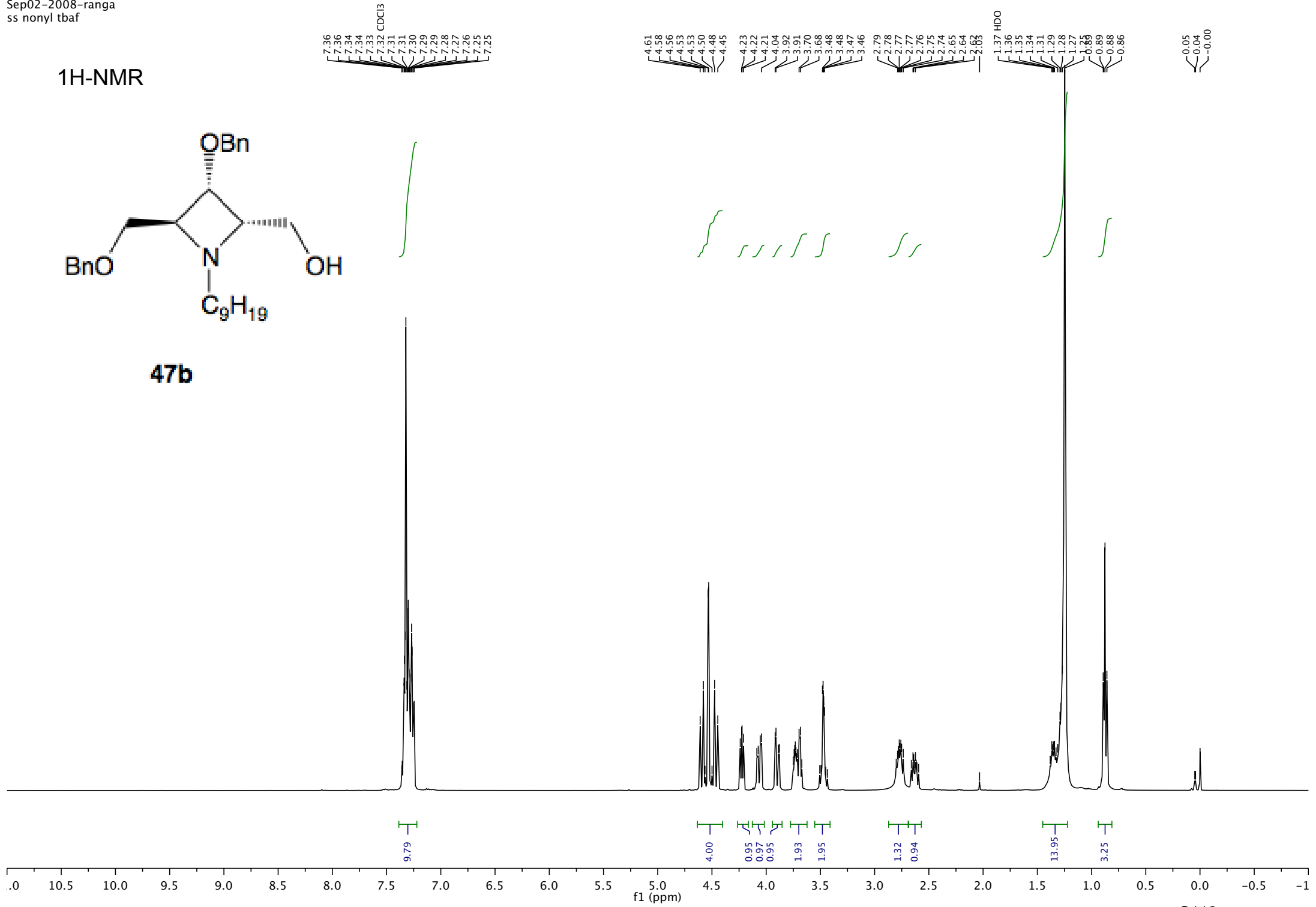
-0.00



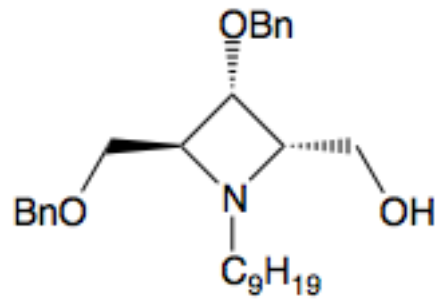
1H-NMR



47b



13C-NMR



47b

138.22
137.64
128.49
128.37
127.88
127.70
127.68
127.63

99.99

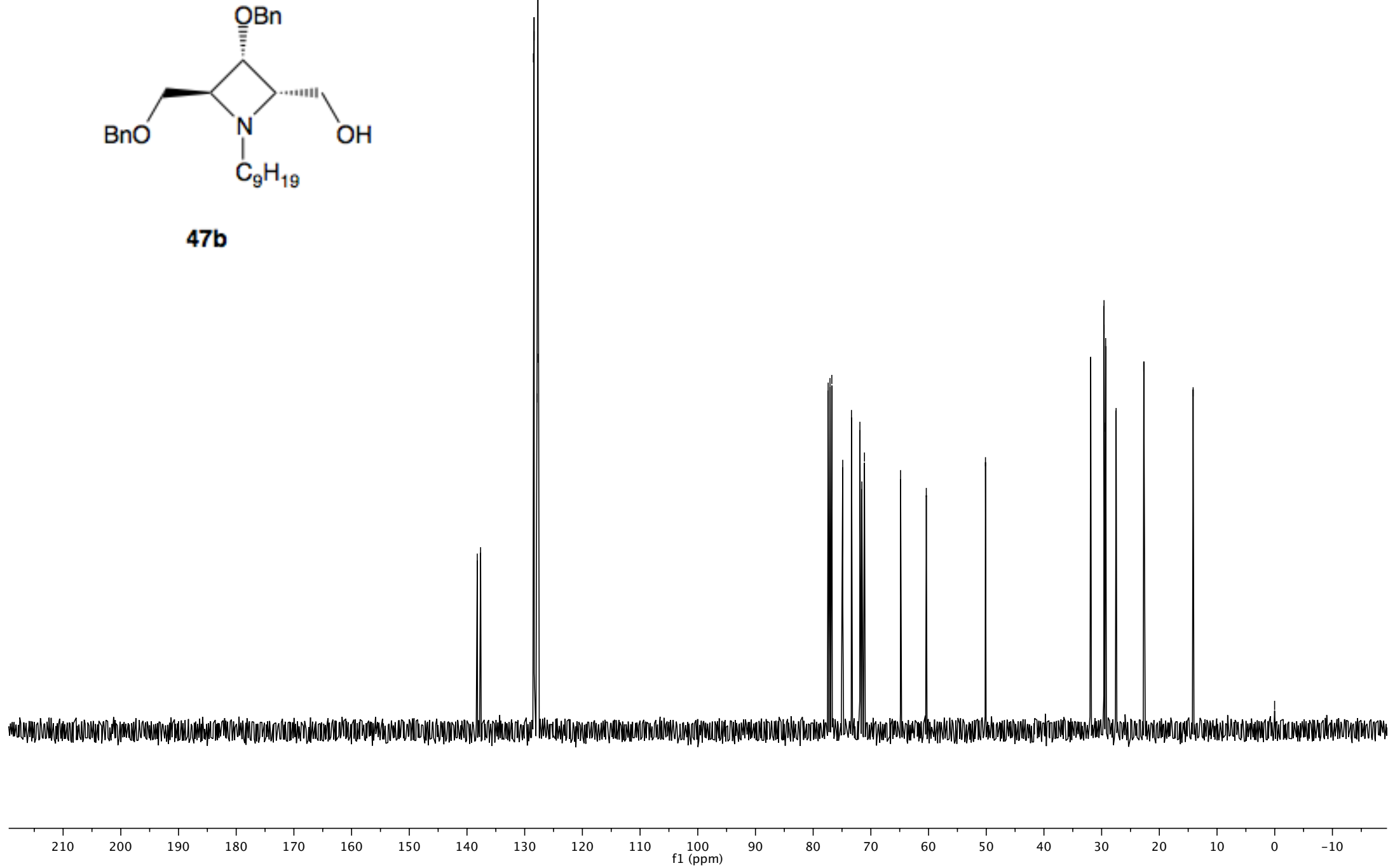
77.40 CDCl₃
77.08 CDCl₃
76.76 CDCl₃
74.88
73.36
71.90
71.56
71.11
64.85
60.39

50.14

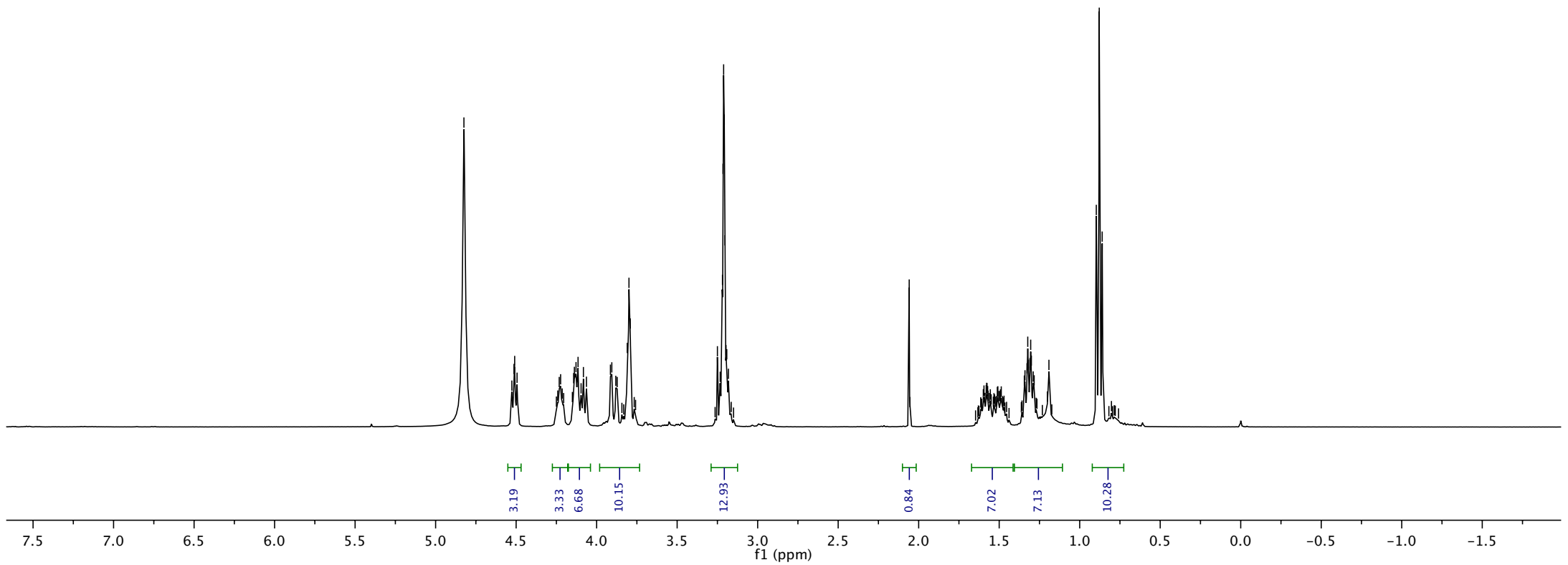
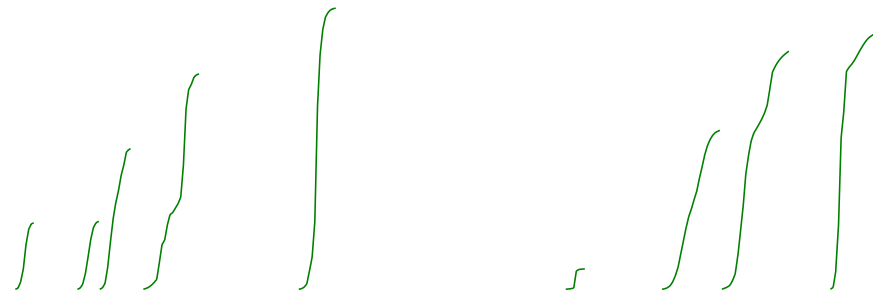
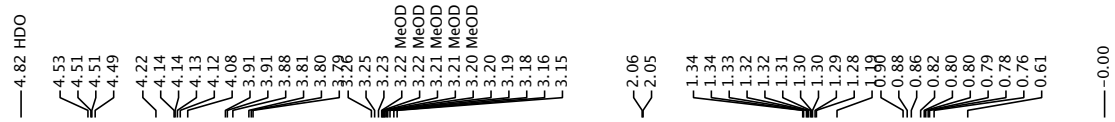
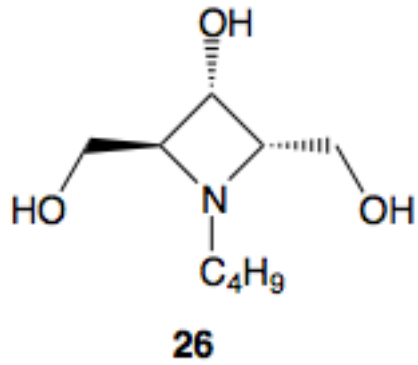
31.90
28.59
28.57
29.34
29.30
27.49
22.70

14.14

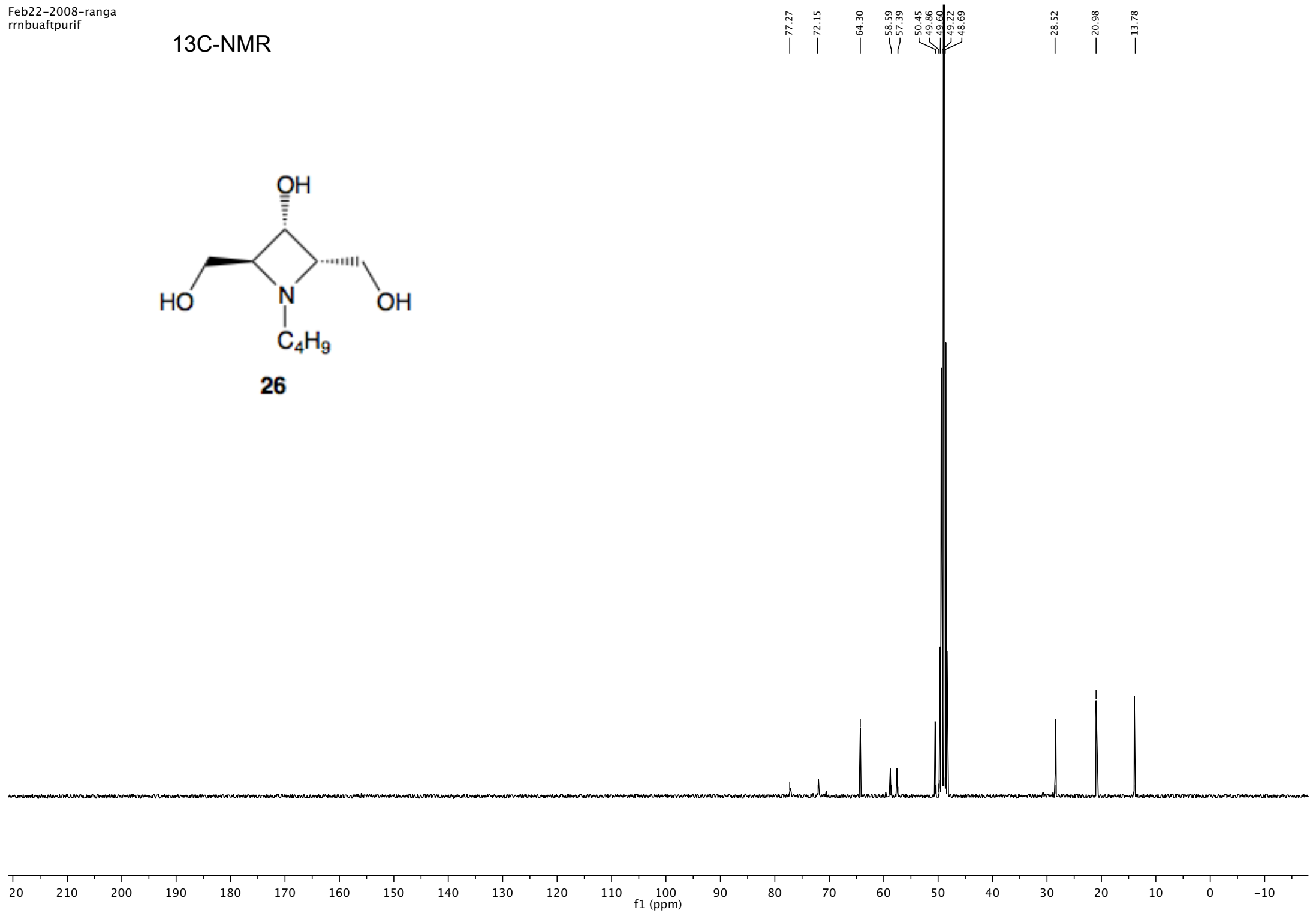
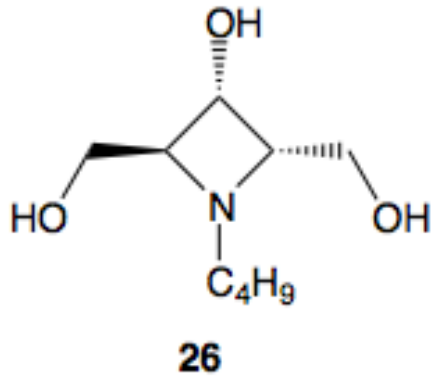
0.02



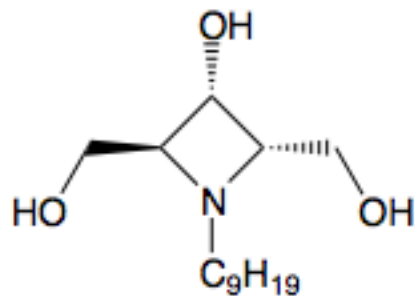
1H-NMR



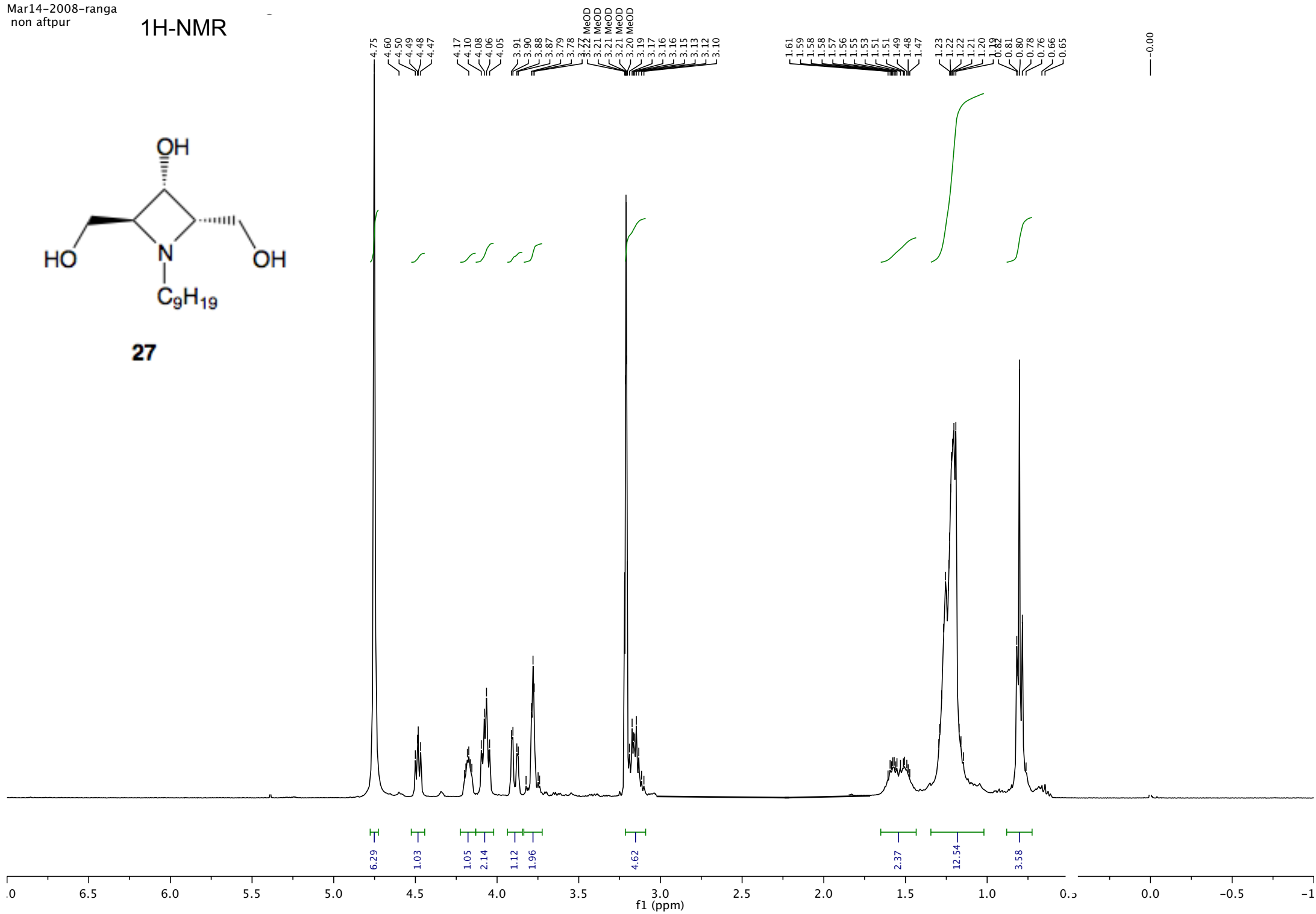
13C-NMR



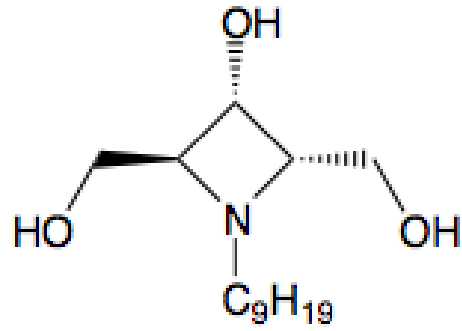
1H-NMR



27



13C-NMR



27

