

Synthesis and cytotoxicity of dibenzo[(γ -aryl)pyridino]aza-17-crown-5 ethers

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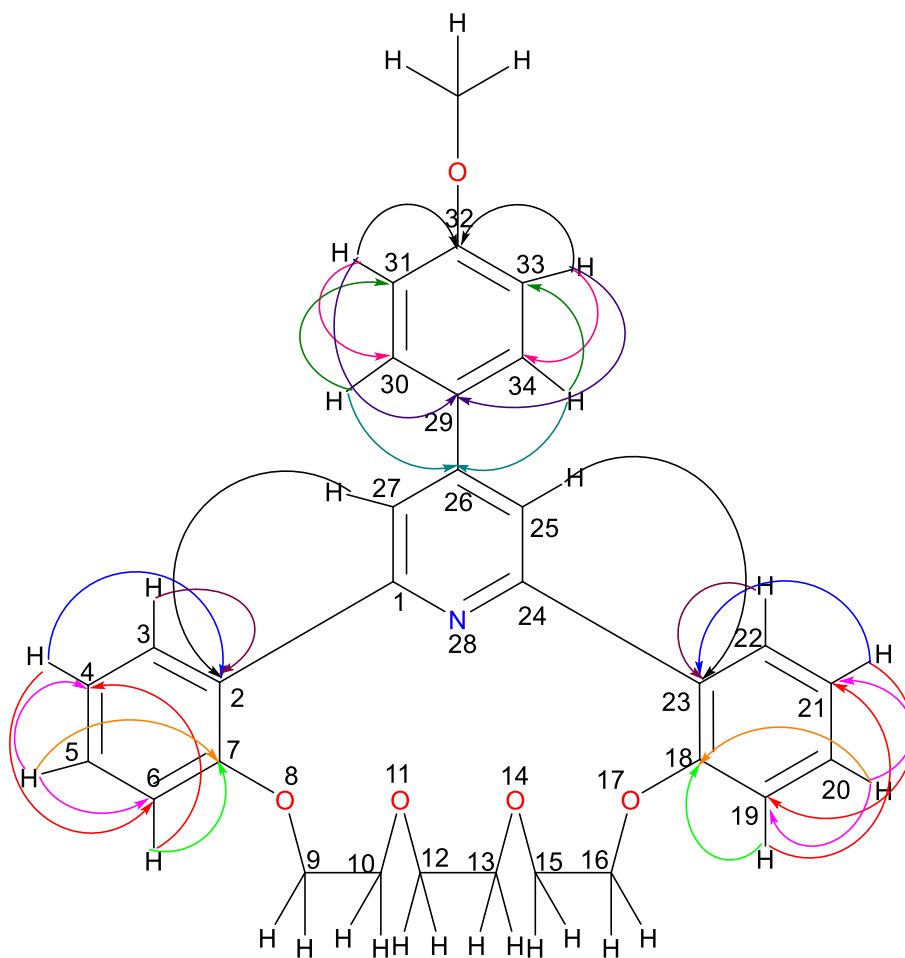
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Result from HMBC and HSQC



N ₀	Positions	¹ H-NMR (ppm)	¹³ C-NMR
1.	12,13	3.19	70.58
2.	10,15	3.62	69.56
3.	OCH ₃	3.89	55.51
4.	9,16	4.10	71.16
5.	4,21	6.98-7.02	114.52
6.	6,19	6.98-7.02	114.62
7.	5,20	7.12	121.63
8.	3,22	7.35	129.67
9.	30,34	7.64-7.68	128.47
10.	31,33	7.64-7.68	130.76
11.	25, 27	7.72	121.95

12.	2, 23	-	131.65
13.	7,18	-	132.60
14.	29	-	146.77
15.	26	-	156.40
16.	1, 24	-	156.94
17.	32	-	160.43

The HMBC (Heteronuclear Multiple Bond Correlation) experiment gives correlations between carbons and protons that are separated by two, three bonds.

H4	C6
H21	C19
H6	C4
H19	C21
H4	C2
H21	C23
H6	C7
H19	C18
H5	C4,5
H20	C19,21
H5	C7
H20	C18
H3	C2
H22	C23
H31	C30
H33	C34
H30	C31
H34	C33
H31	C29
H33	C29
H30	C26
H34	C26
H31	C32

H33	C32
H27	C2
H25	C23

In aromatic zone showed cross peaks and the assignment of their carbon resonances: H-4 and H-21 (σ H 6.98-7.02 $\rightarrow$ C-2 and C-23, respectively (σ C 131.65), C-6 and C-19 (σ C 114.62); H-6 and H-19 (σ H 6.98-7.02) $\rightarrow$ C-4 and C-21 (σ C 114.52), C-7 and C-18 (σ C 132.60); H-5 and H-20 (σ H 7.12) $\rightarrow$ C-4 and C-21 (σ C 114.52), C-6 and C19 (σ C 114.62) and C-7 and C-18 (σ C 132.60); H-3 and H-22 (σ H 7.35) $\rightarrow$ C2 and C23 (σ C 131.65), H-31 and H-33 (σ H 7.64-7.68) $\rightarrow$ C30 and C-34 (σ C 128.47), C-29 (σ C 146.77), C-32 (σ C 160.43), -30 and H-34 $\rightarrow$ C-26 (σ C 156.40), C31 and C-33 (σ C 130.76), H-25 and H-27 (σ H 7.72) $\rightarrow$ C-2 and C-23, respectively (σ C 131.65).

X-ray diffraction experiment

The crystal of **3a** ($\text{C}_{30}\text{H}_{29}\text{NO}_5$, $M = 483.54$) is monoclinic, space group $P2_1/c$, at $T = 120$ K: $a = 10.239(4)$ Å, $b = 10.344(4)$ Å, $c = 23.413(8)$ Å, $\beta = 98.648(6)^\circ$, $V = 2451.5(15)$ Å³, $Z = 4$, $d_{\text{calc}} = 1.310$ g/cm³, $F(000) = 1024$, $\mu = 0.089$ mm⁻¹. 21314 total reflections (5116 unique reflections, $R_{\text{int}} = 0.082$) were measured on a three-circle Bruker APEX-II CCD diffractometer ($\lambda(\text{MoK}\alpha)$ -radiation, graphite monochromator, φ and ω scan mode, $2\theta_{\text{max}} = 54.0^\circ$) and corrected for absorption ($T_{\text{min}} = 0.975$; $T_{\text{max}} = 0.980$).^[1] The structure was determined by direct methods and refined by full-matrix least squares technique on F^2 with anisotropic displacement parameters for non-hydrogen atoms. The hydrogen atoms were placed in calculated positions and refined within riding model with fixed isotropic displacement parameters [$U_{\text{iso}}(\text{H}) = 1.5U_{\text{eq}}(\text{C})$ for the methyl group and $1.2U_{\text{eq}}(\text{C})$ for the other groups]. The final divergence factors were $R_1 = 0.116$ for 3809 independent reflections with $I > 2\sigma(I)$ and $wR_2 = 0.242$ for all independent reflections, $S = 1.031$. All calculations were carried out using the SHELXTL program.^[2]

Crystallographic data for **3a** have been deposited with the Cambridge Crystallographic Data Center, CCDC 1532287. Copies of this information may be obtained free of charge from the Director, CCDC, 12 Union Road, Cambridge CB2 1EZ, UK (fax: +44 1223 336033; e-mail: deposit@ccdc.cam.ac.uk or www.ccdc.cam.ac.uk).

Table 1. Hydrogen bonds for **3a** [Å and °].

$\text{D}^*-\text{H}\cdots\text{A}^*$	$d(\text{D}-\text{H})$	$d(\text{H}\cdots\text{A})$	$d(\text{D}\cdots\text{A})$	$\angle\text{DHA}$
$\text{C9}-\text{H9a}\cdots\text{N28}^a$	0.99	2.59	3.491(7)	151.9
$\text{C12}-\text{H12a}\cdots\text{O17}^b$	0.99	2.56	3.335(7)	135.1
$\text{C12}-\text{H12b}\cdots\text{O8}$	0.99	2.39	3.013(6)	119.9
$\text{C13}-\text{H13b}\cdots\text{O17}$	0.99	2.55	2.958(6)	104.8
$\text{C35}-\text{H35a}\cdots\text{O11}^c$	0.98	2.55	3.465(7)	156.1

* D – proton donor; A – proton acceptor;

Symmetry transformations used to generate equivalent atoms:

^a $-x, -y+1, -z$; ^b $-x, y+0.5, -z+0.5$; ^c $x+1, -y+1.5, z+0.5$

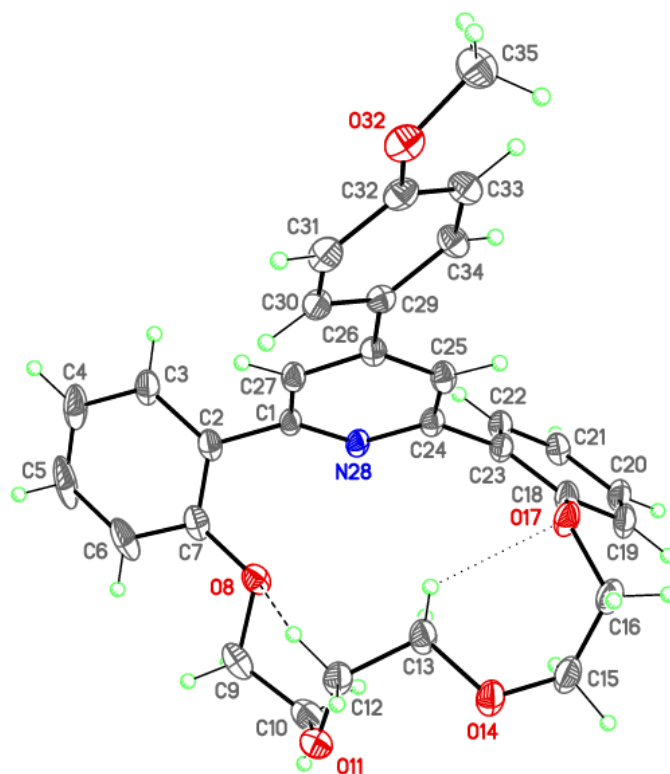


Figure 1. Molecular structure of **3a**. The dashed and dotted lines indicate the intramolecular C—H···O hydrogen bonds.

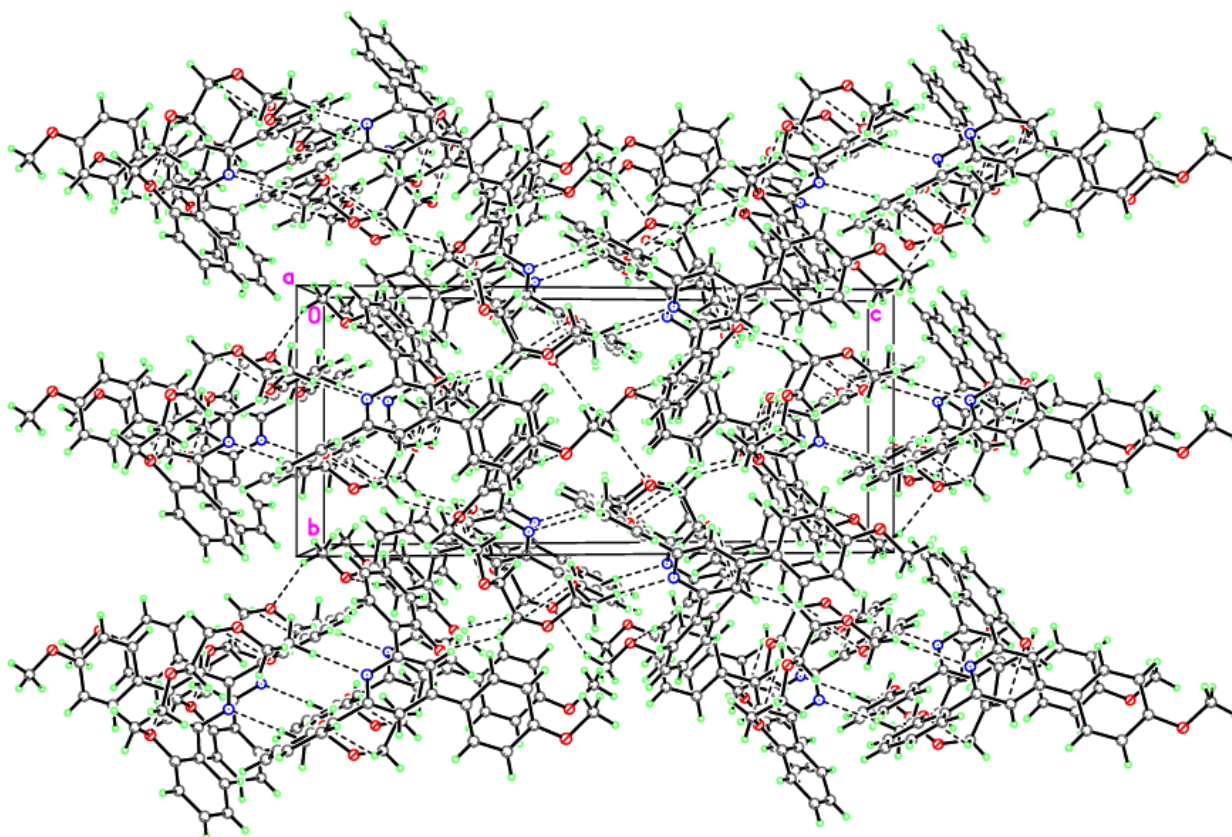


Figure 2. Crystal packing of **3a**. The dashed and dotted lines indicate the intramolecular C—H···O and intermolecular C—H···N and C—H···O hydrogen bonds.

Cytotoxicity assay

Four human cancer cell lines was obtained from the American Type Culture Collection (Manassas, VA) ATCC as FL (human cervix carcinoma), RD (human rhabdosarcoma), Lu1 (human lung adenocarcinoma), HepG2 (human hepatocellular carcinoma). The Vero cell line was initiated from kidney of a normal adult African green monkey. They were grown in DMEM (Dulbecco's Modified Eagle Medium) (Thermo Fisher Scientific, USA) supplemented with L-glutamine, Sodium pyruvate, NaHCO₃, PSF (Penicillin- Streptomycin sulfate - Fungizone); NAA (Non-Essential Amino Acids); 10% BCS (Bovine Calf Serum). All incubations were performed at 37°C for 72 hours in a CO₂ (5%) incubator with the plates capped in the normal fashion

The MTT is based on the protocol described by Skehan & CS (1990) and Likhiwitayawuid & CS (1993).^[3] This method has worldwide application and is recommended by National Cancer Institute (NCI) and College of Medicine, University of Illinois at Chicago for routine drug screening.

Table 1. Cytotoxicity tests performed on compounds **3 a-g** in human cancer cell lines.

№	Comp.	Conc. (µg/ml)	Cell line, cell survival (%)				Concl.
			HepG2	Lu1	RD	FL	
	DMSO	-	100	100	100	100	
	Taxol (+)	5	1,34±0,8	3,66±0,9	0,68±0,3	3,87±1,1	Positive
1	3 a	10	100	99.06±1.5	100	100	Negative
2	3 b	10	0	0	0	0	Positive with 04 cell lines
3	3 c	10	0	0	0	0	Positive with 04 cell lines
4	3 d	10	20.07±0.8	38.25±0.4	39.26±0.7	64.09±0.8	Positive with 03 cell lines

5	3 e	10	67,61±1,4	64,44±1,5	83,27±1,2	59,3±2,3	Negative
6	3 f	10	58,45±0,8	72,89±2,8	44,83±1,1	13,46±1,6	Positive with 02 cell lines: RD, FL
7	3 g	10	60,85±1,4	69,47±2,6	50,09±2,8	41,71±3,1	Positive with 02 cell lines: RD, FL

Table 2. Results of IC₅₀ test

Entry	Comp.	Cell lines IC ₅₀ (µg/ml)				Conclusion
		HepG2	Lu1	RD	FL	
	Taxol (+)	0.29	0.31	0.25	0.33	Positive
1	3 b	2.607	2.664	2.56	1.392	Positive with 04 cell lines
2	3 c	7.957	6.946	6.887	6.742	Positive with 04 cell lines
3	3 d	6.586	7.904	7.528	-	Positive with 03 cell lines
4	3 f	-	-	8.881	8.346	Positive with 02 cell lines: RD, FL
5	3 g	-	-	10.0	9.505	Positive with 02 cell lines: RD, FL

Table 3. Results of test on cytotoxicity on the Vero cell

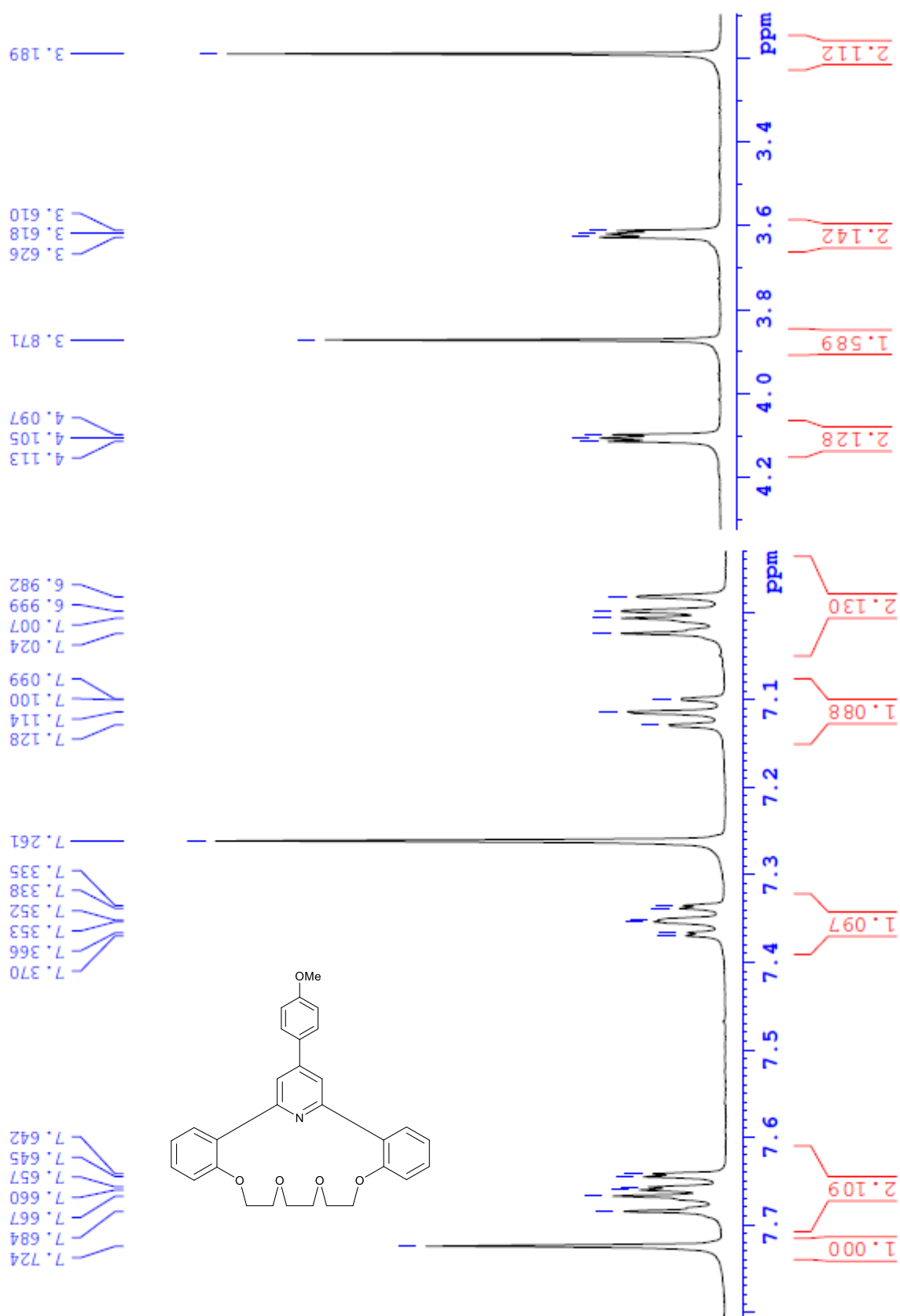
Entry	Compound	Conc. ($\mu\text{g/ml}$)	Vero cell line, cell survival (%)	Conclusion
	DMSO	-	100	
	Taxol (+)	5	32.24$\pm$1.7	
1	3 b	10	71,60 $\pm$ 2,3	Negative
		5	86.55 $\pm$ 0.7	Negative
2	3 c	10	65.86 $\pm$ 0.8	Negative
		5	92.22 $\pm$ 1.5	Negative
3	3 d	10	80.92 $\pm$ 1.3	Negative

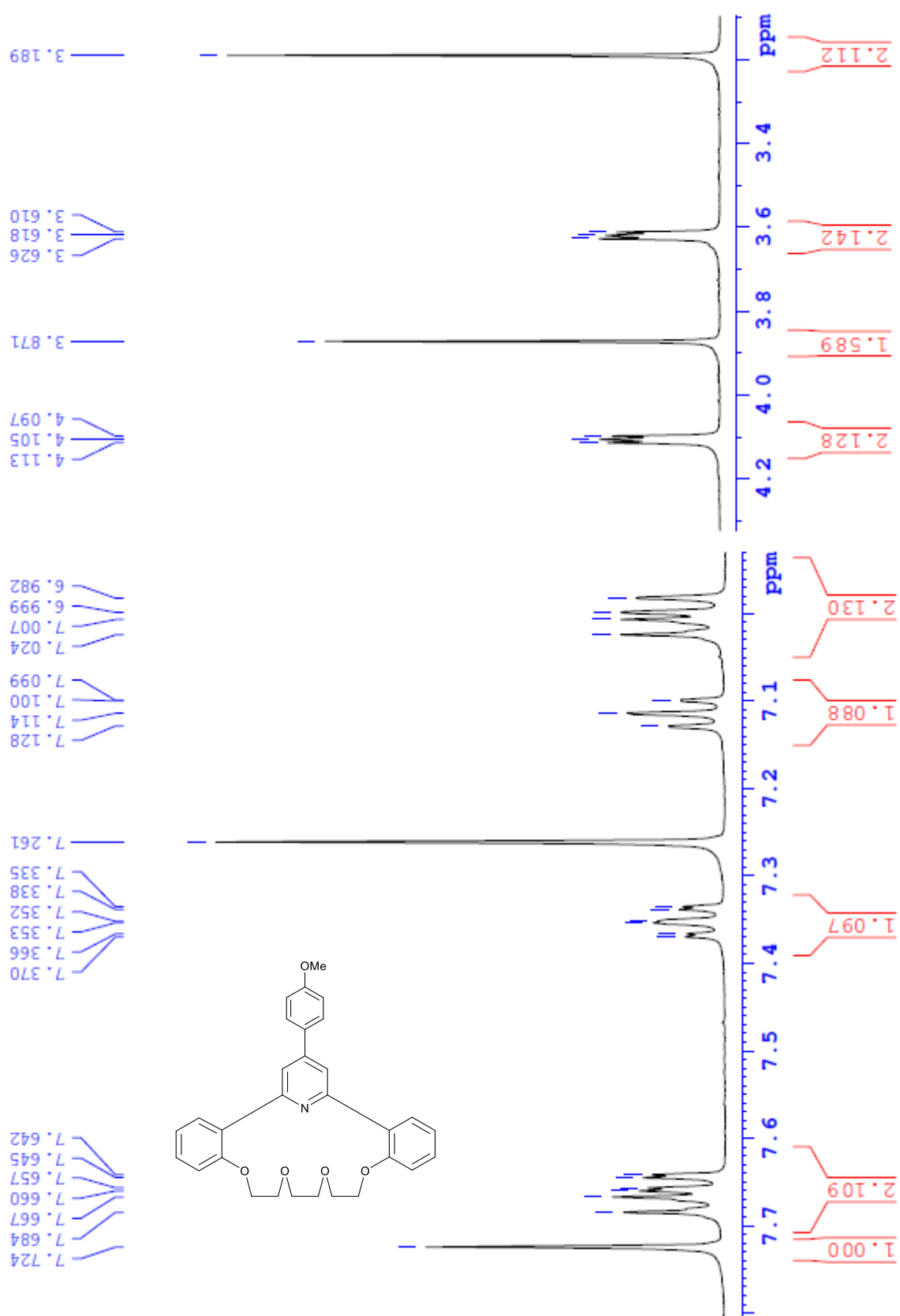
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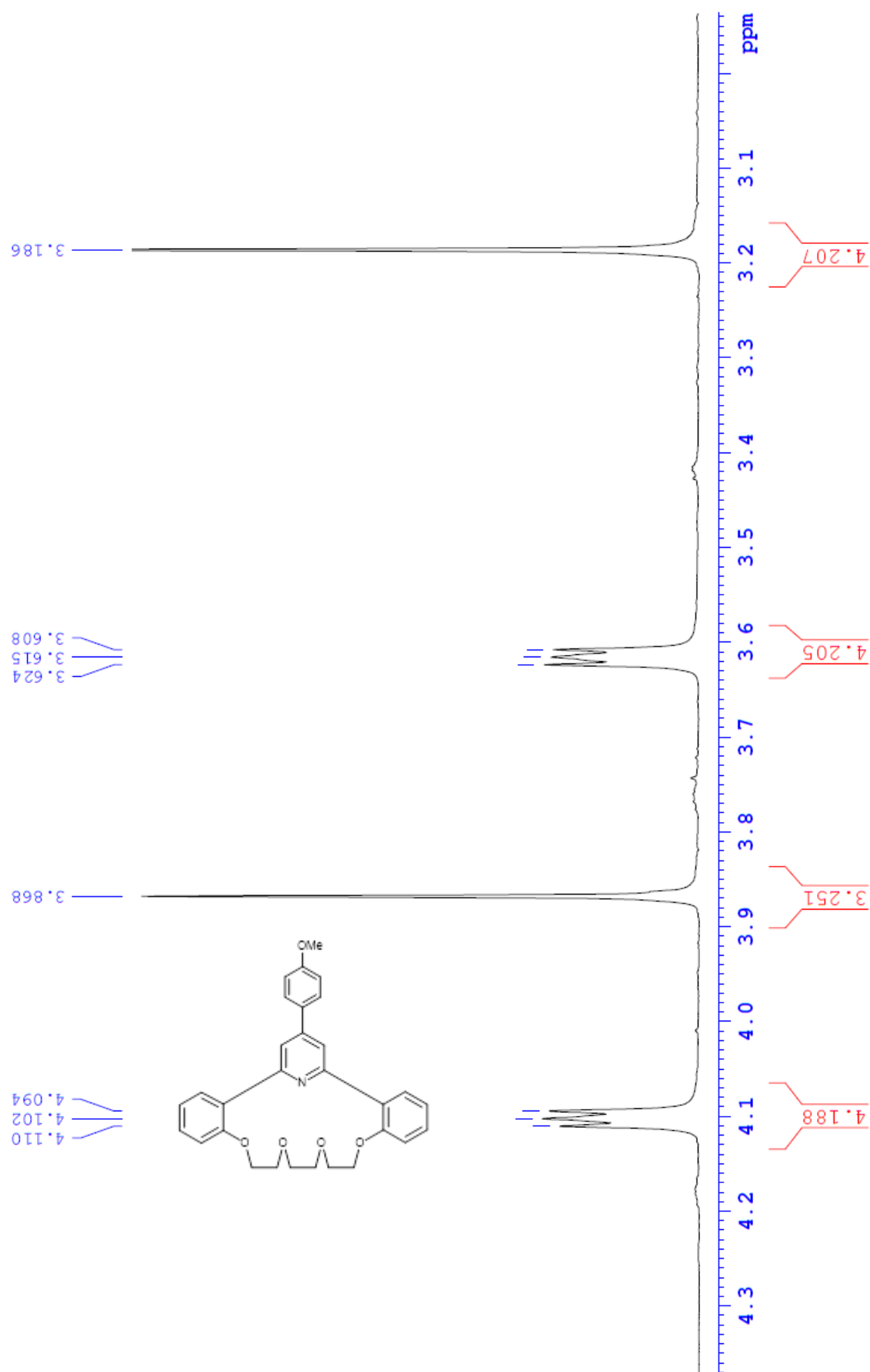
1. G. M. Sheldrick, *SADABS*, v. 2.03, Bruker/Siemens Area Detector Absorption Correction Program; Bruker AXS Inc., Madison, WI, **2003**.
2. G. M. Sheldrick, *Acta Cryst.* **2015**, *C71*, 3-8.
3. Skehan P., Storeng R., Scudiero D., Monks A., McMahon J., Vistica D., Warren J.T., Bokesch H., Kenney S., Boyd M.R. New colorimetric cytotoxicity assay for anticancer-drug screening. *J Natl Cancer Inst.* **1990**, 82(13),1107–1112.
4. Likhitwitayawuid K., Angerhofer C.K., Cordell G.A., Pezzuto J.M., Ruangrunsi N. Cytotoxic and antimalarial bisbenzylisoquinoline alkaloids from *Sephania erecta*. *Journal of Natural Products*. **1993**, 56 (1) pp. 30-38.

Copies of HRMS, ¹H NMR and ¹³C NMR spectra

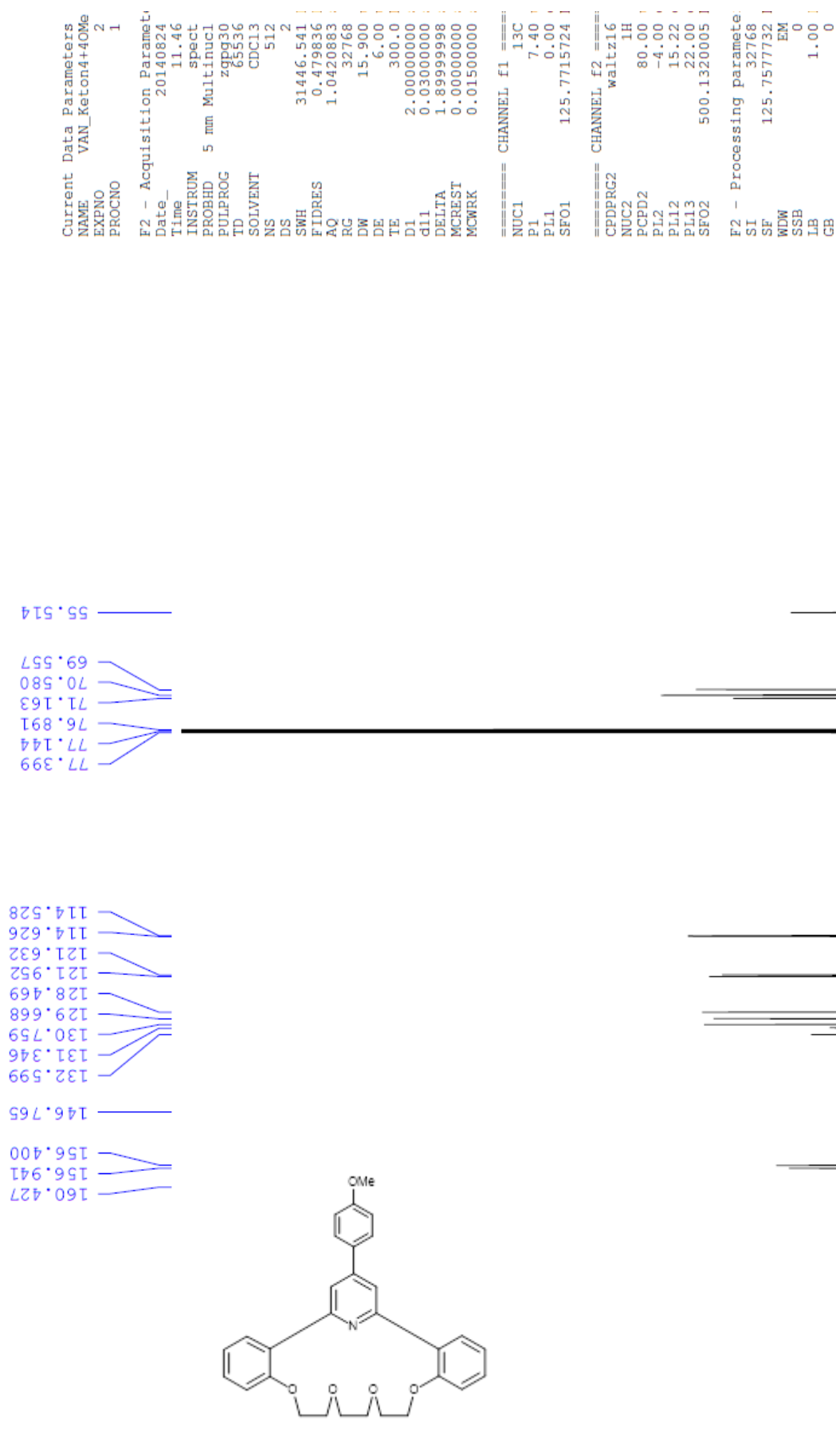
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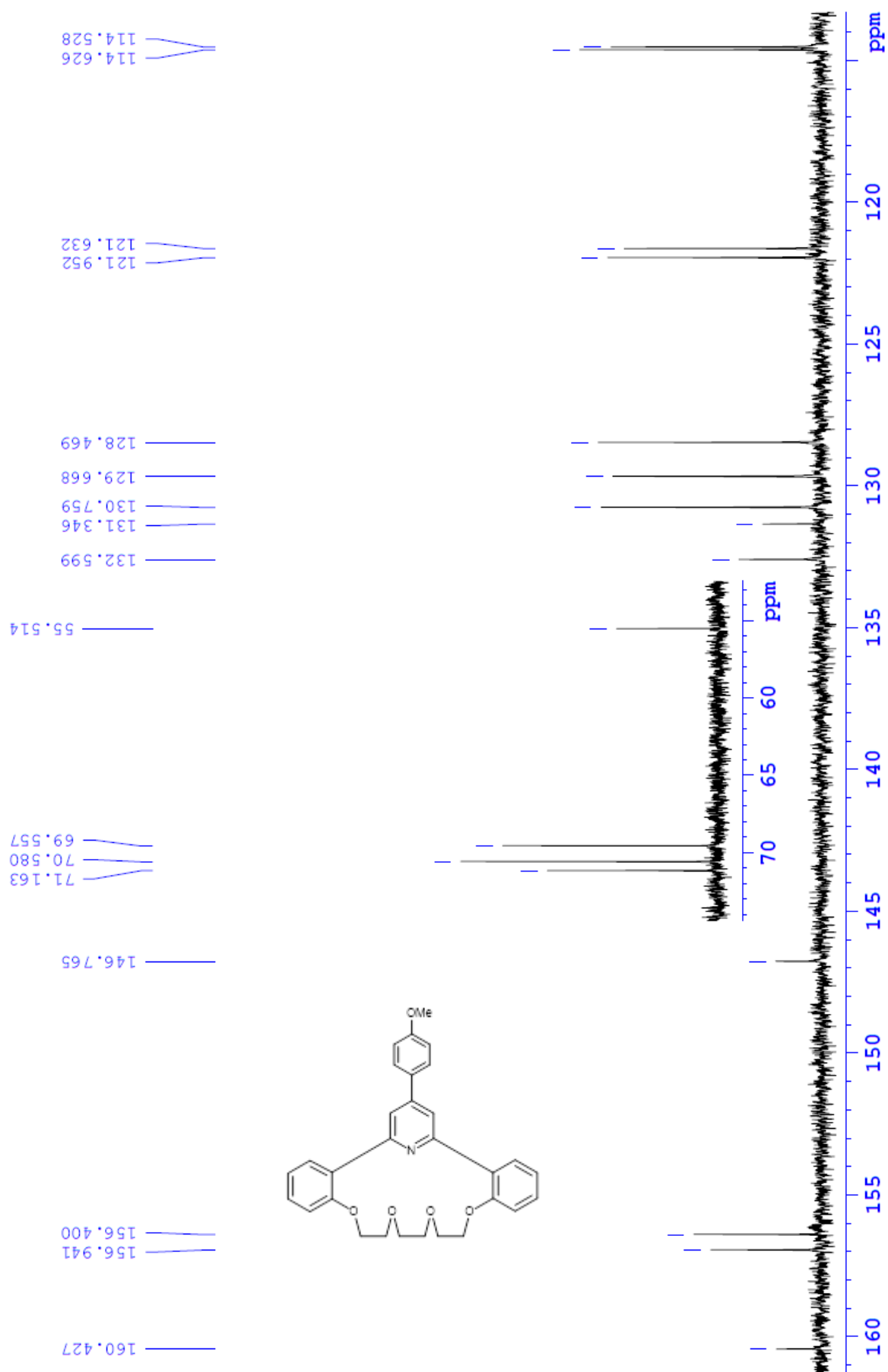


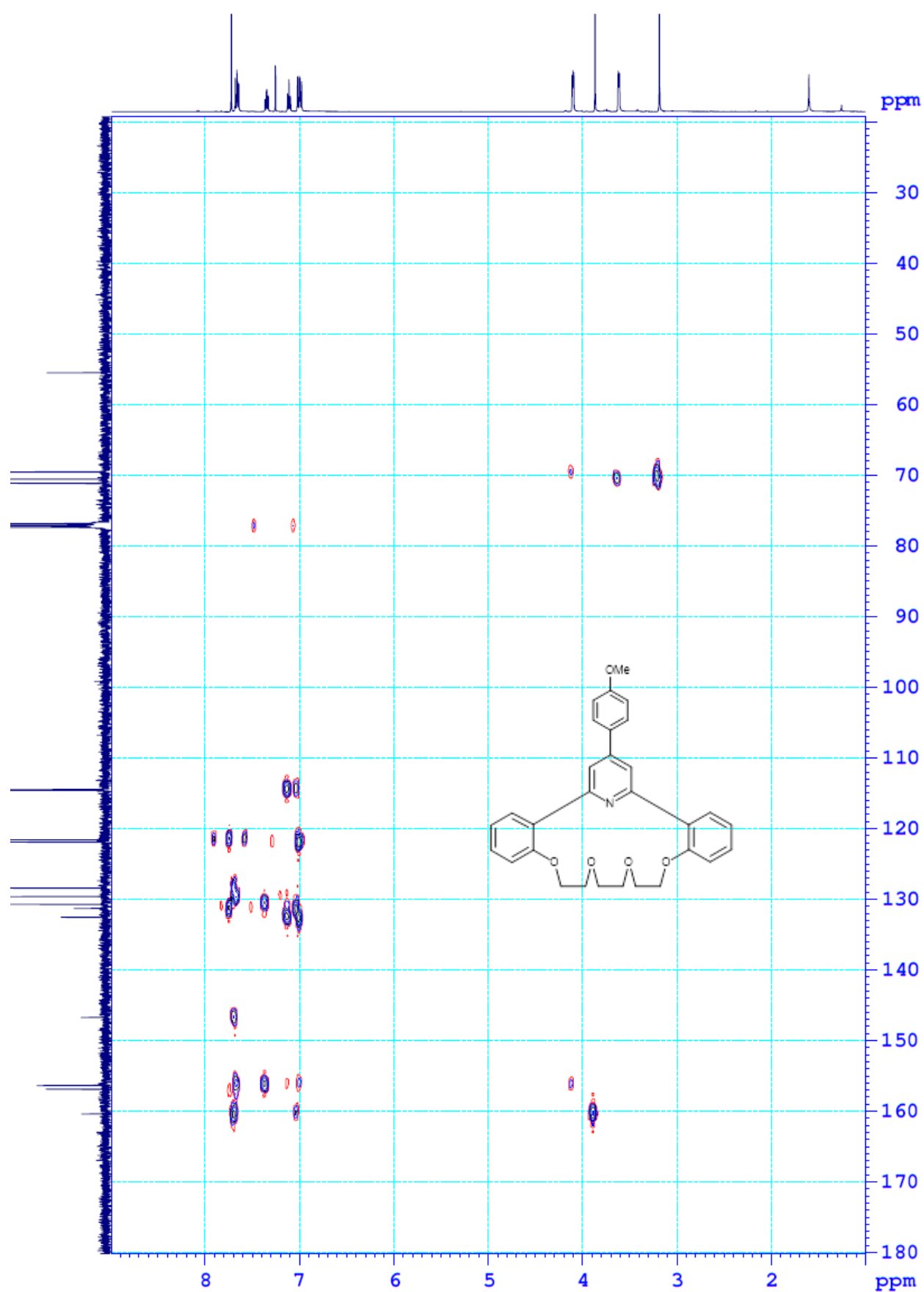


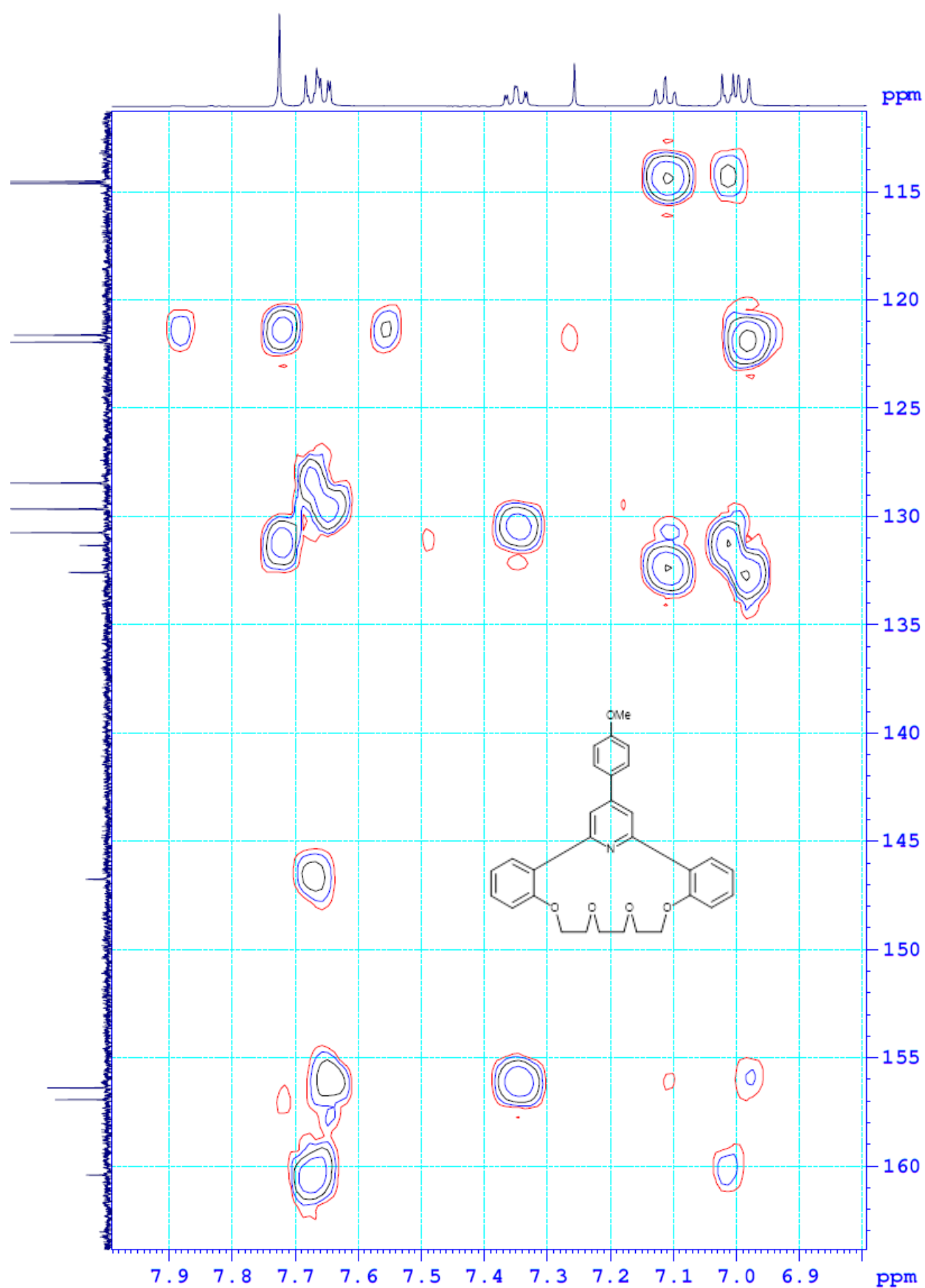


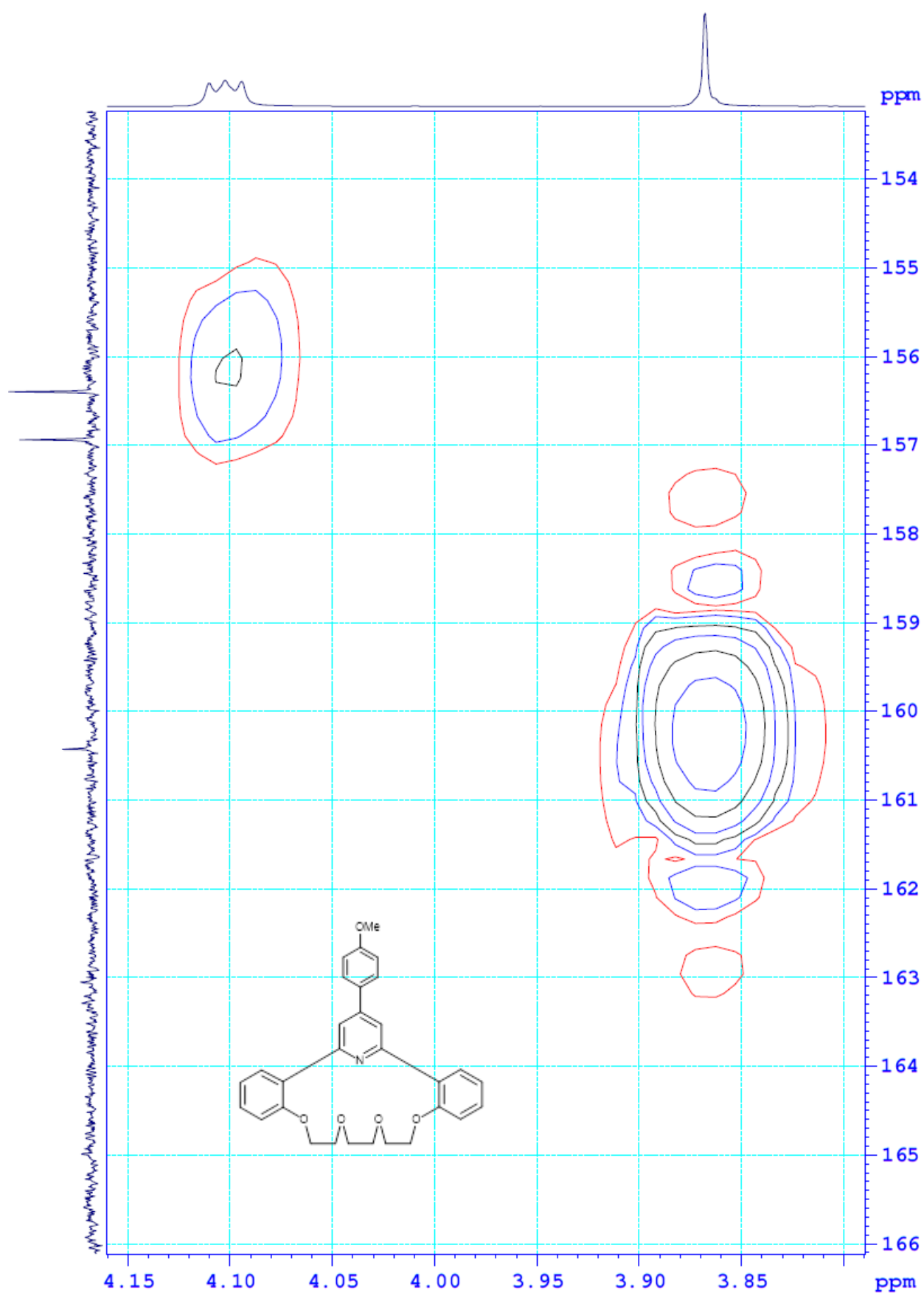
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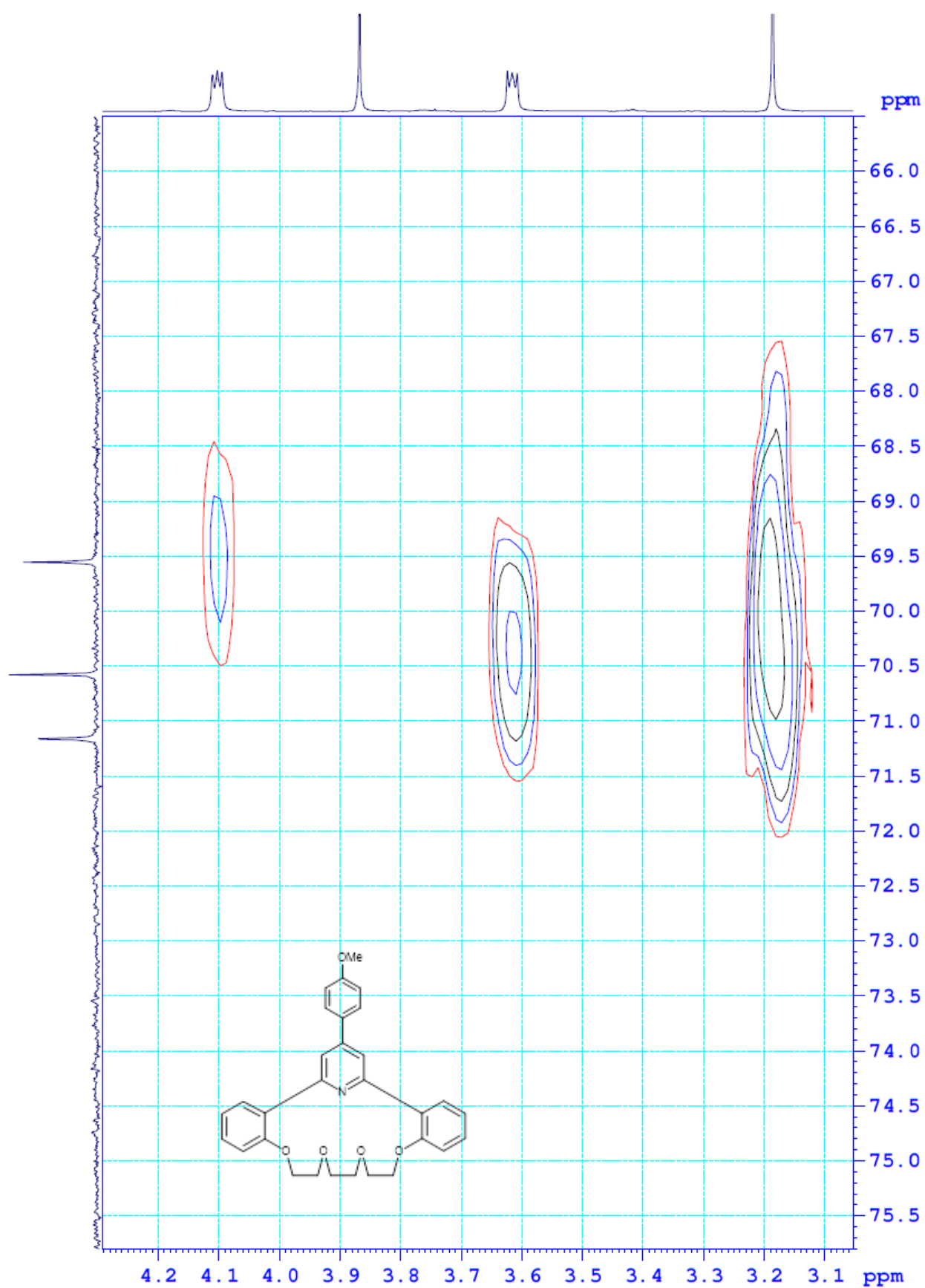


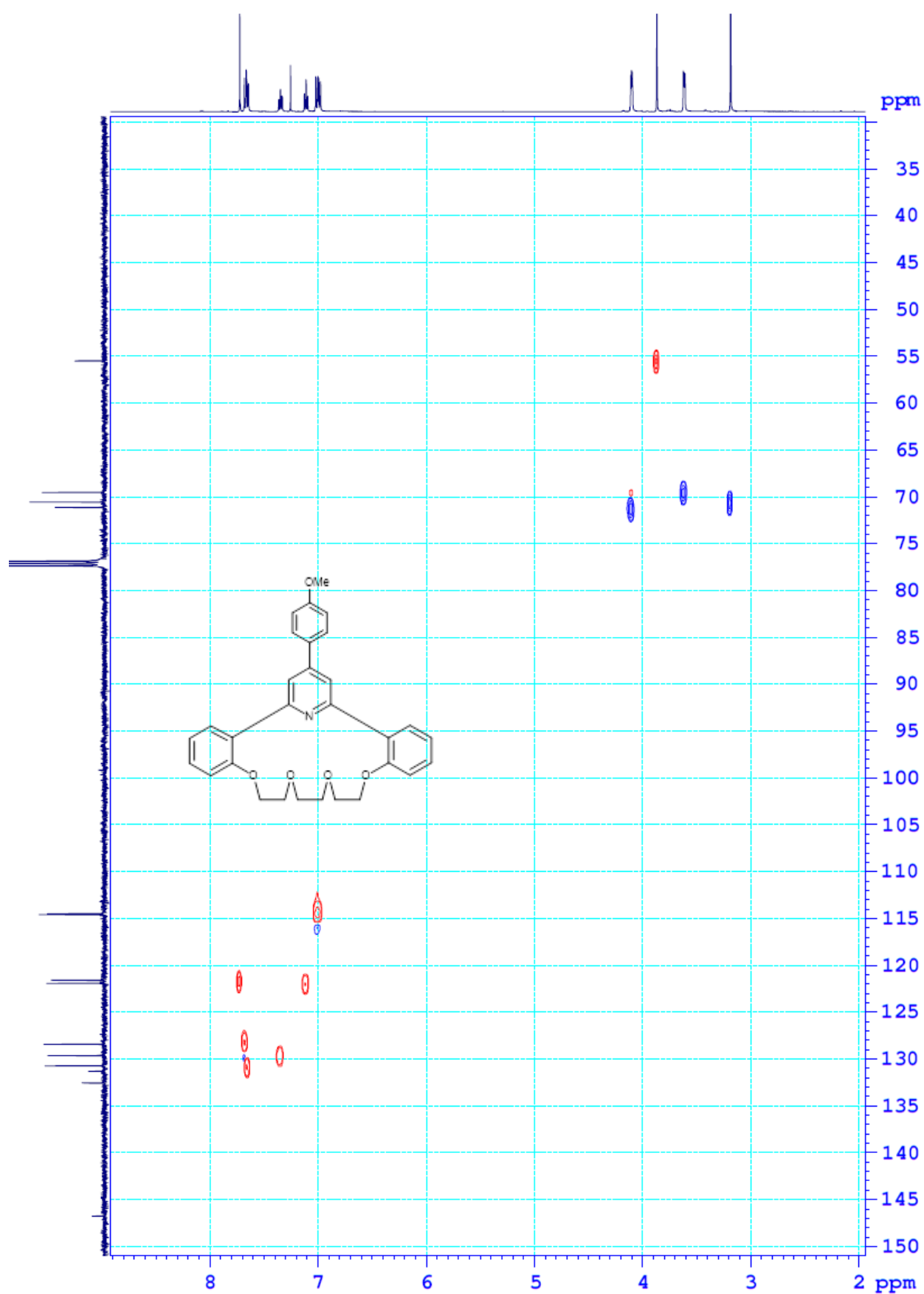
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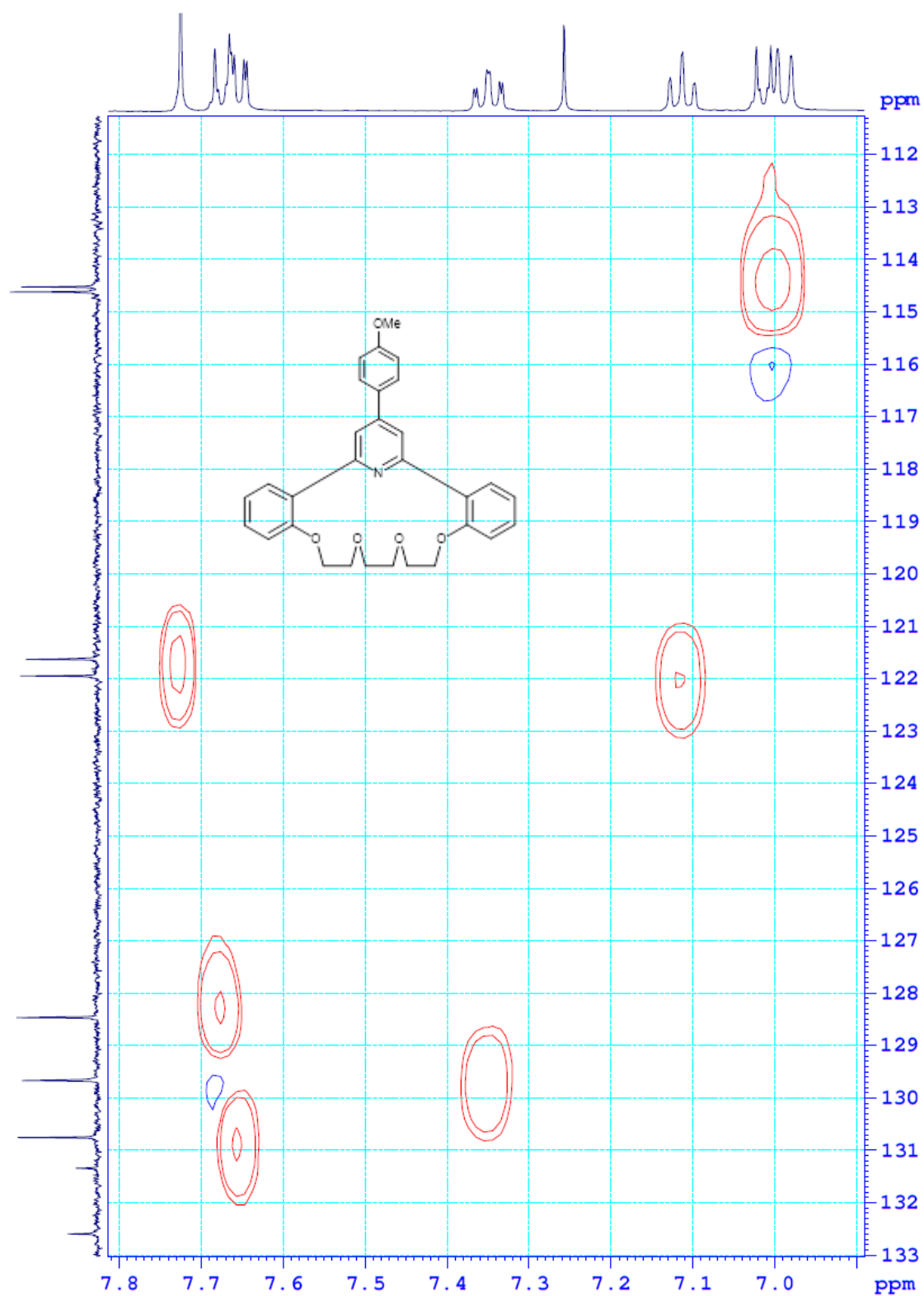
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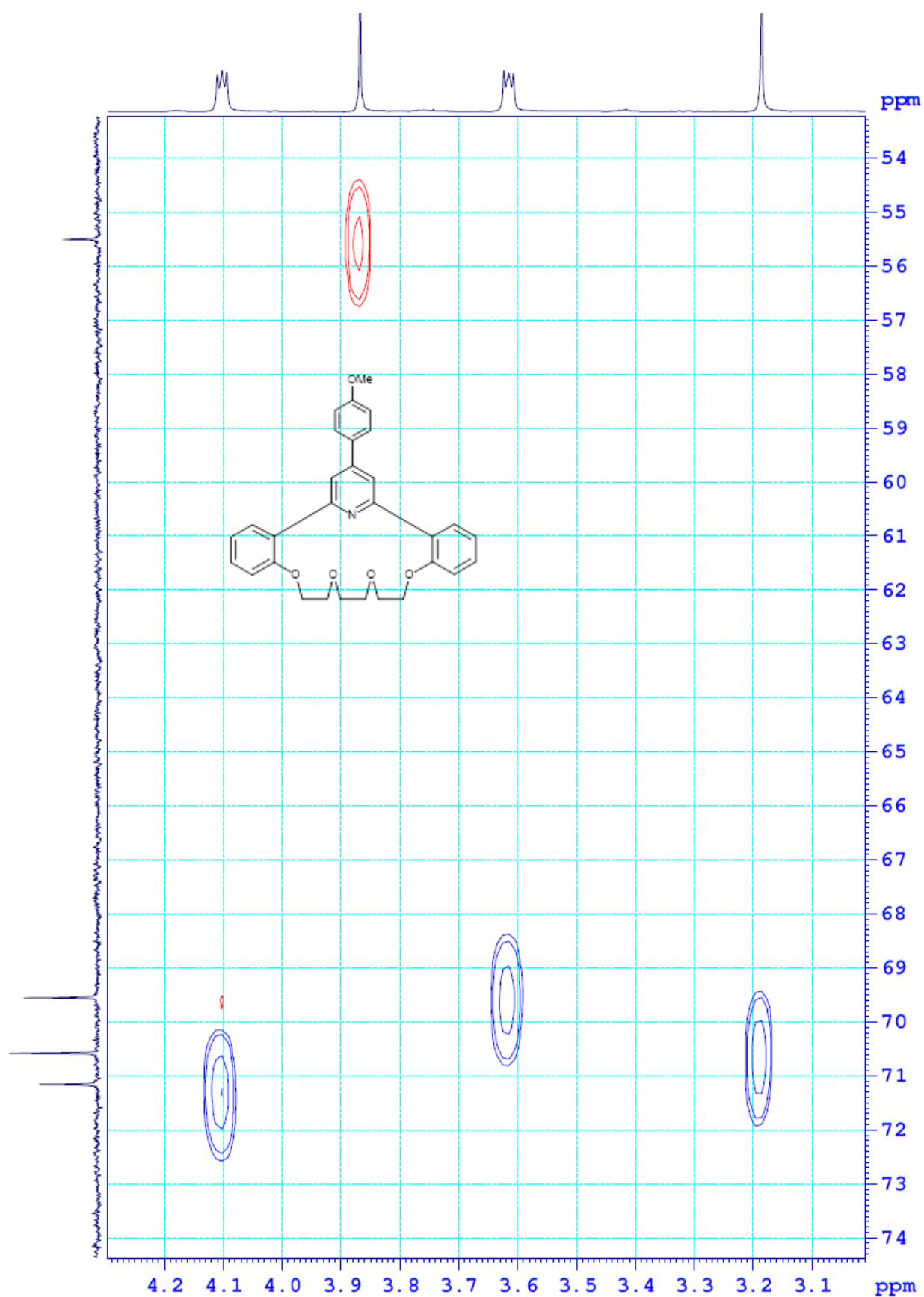
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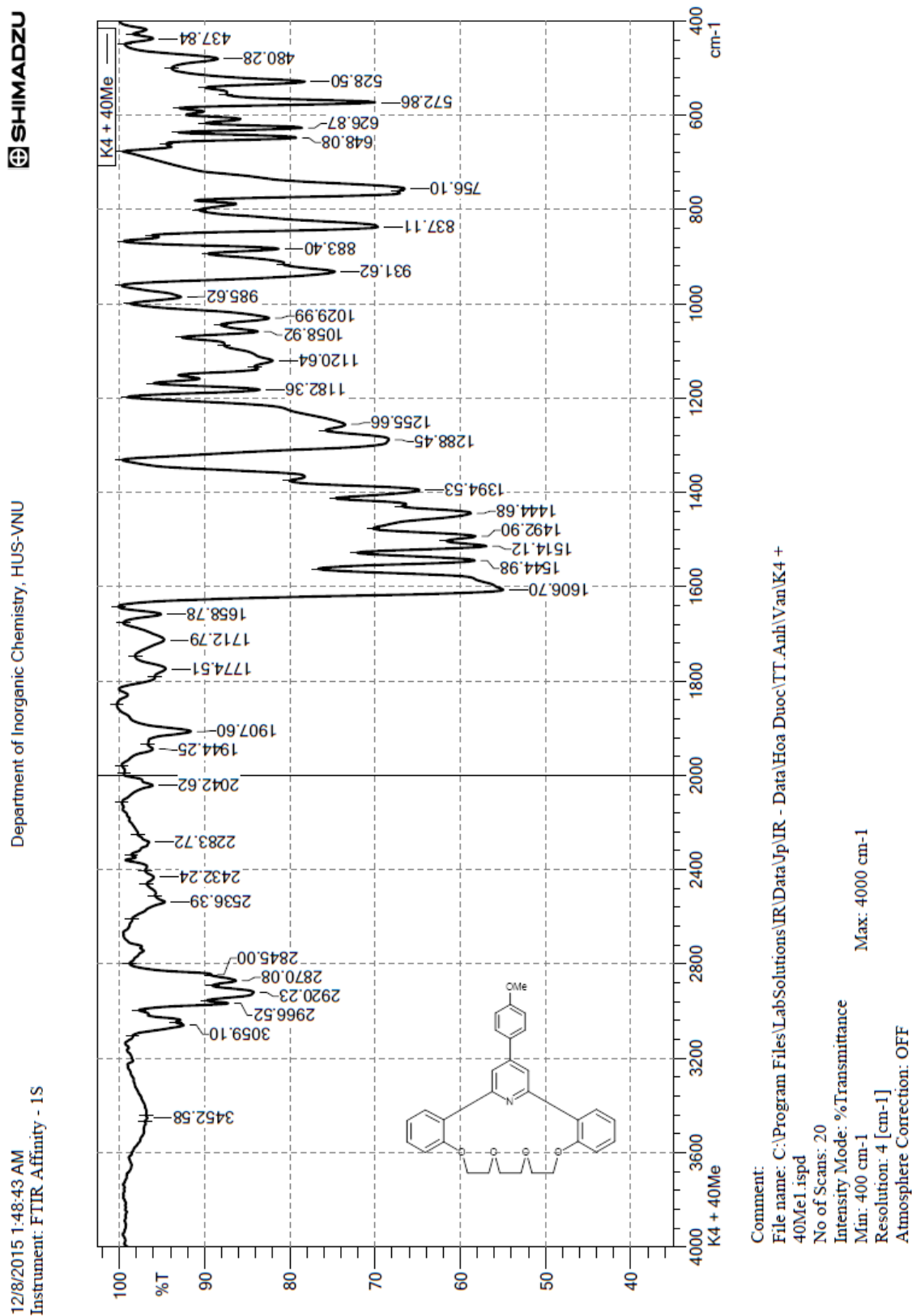
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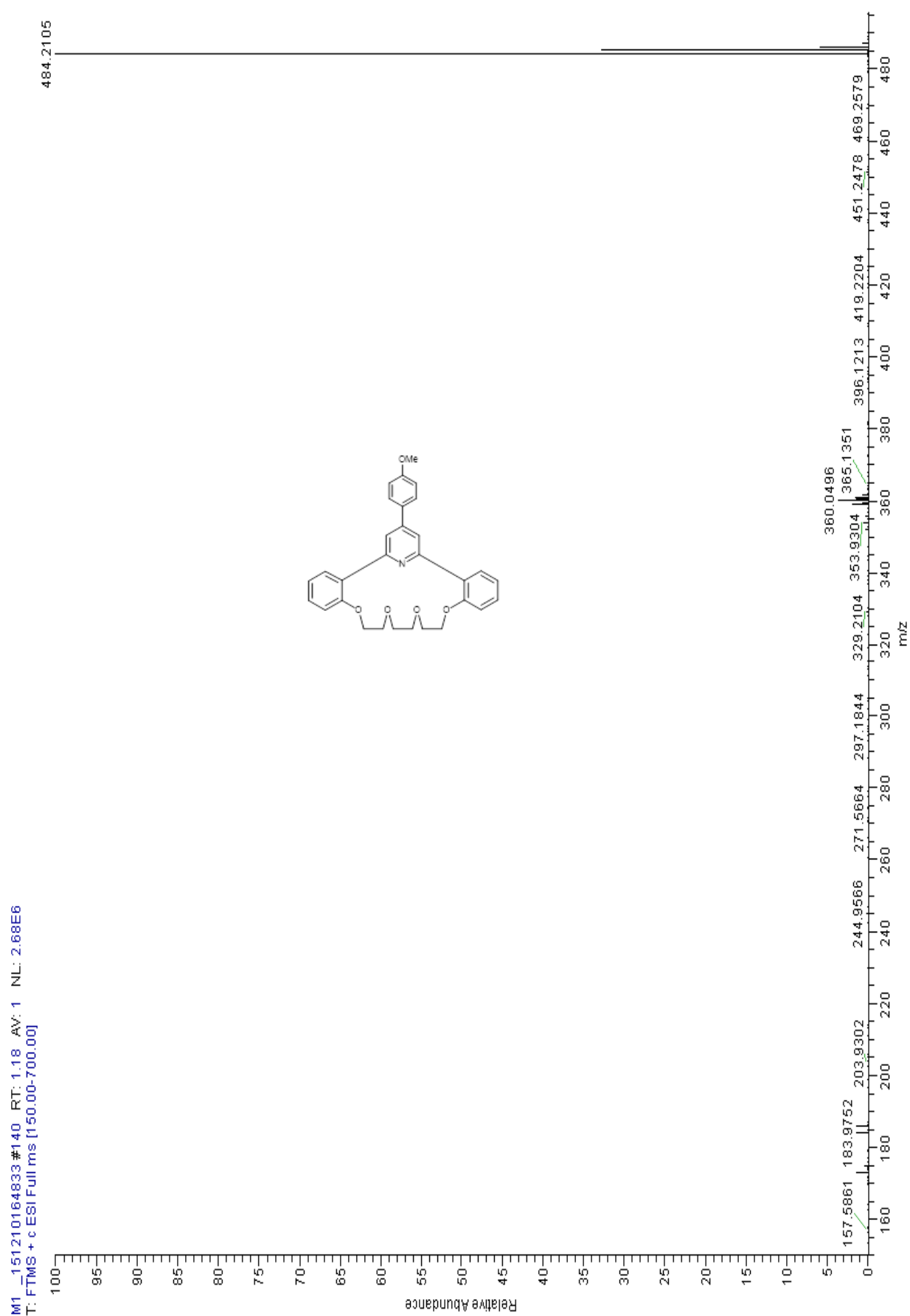


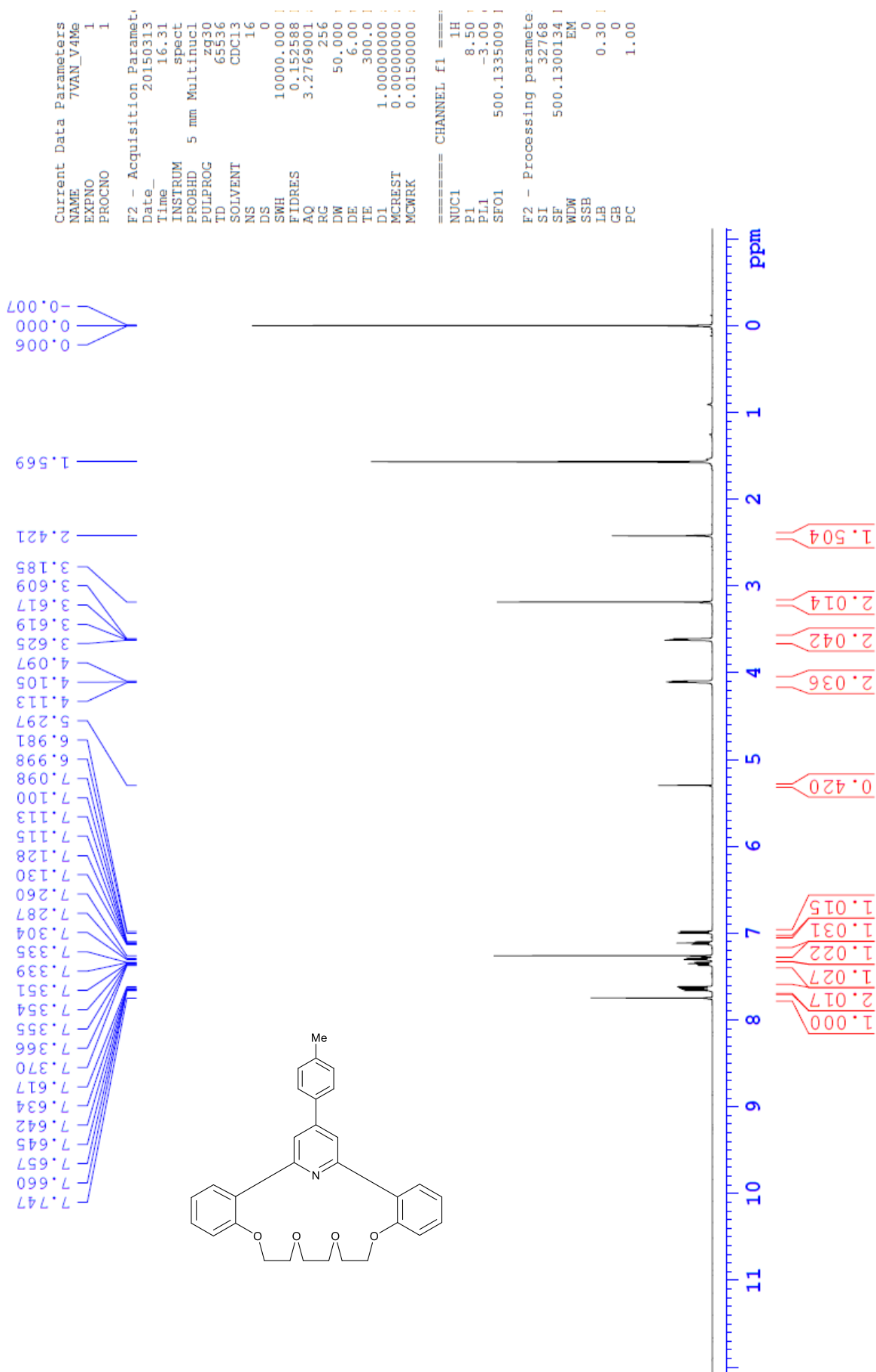
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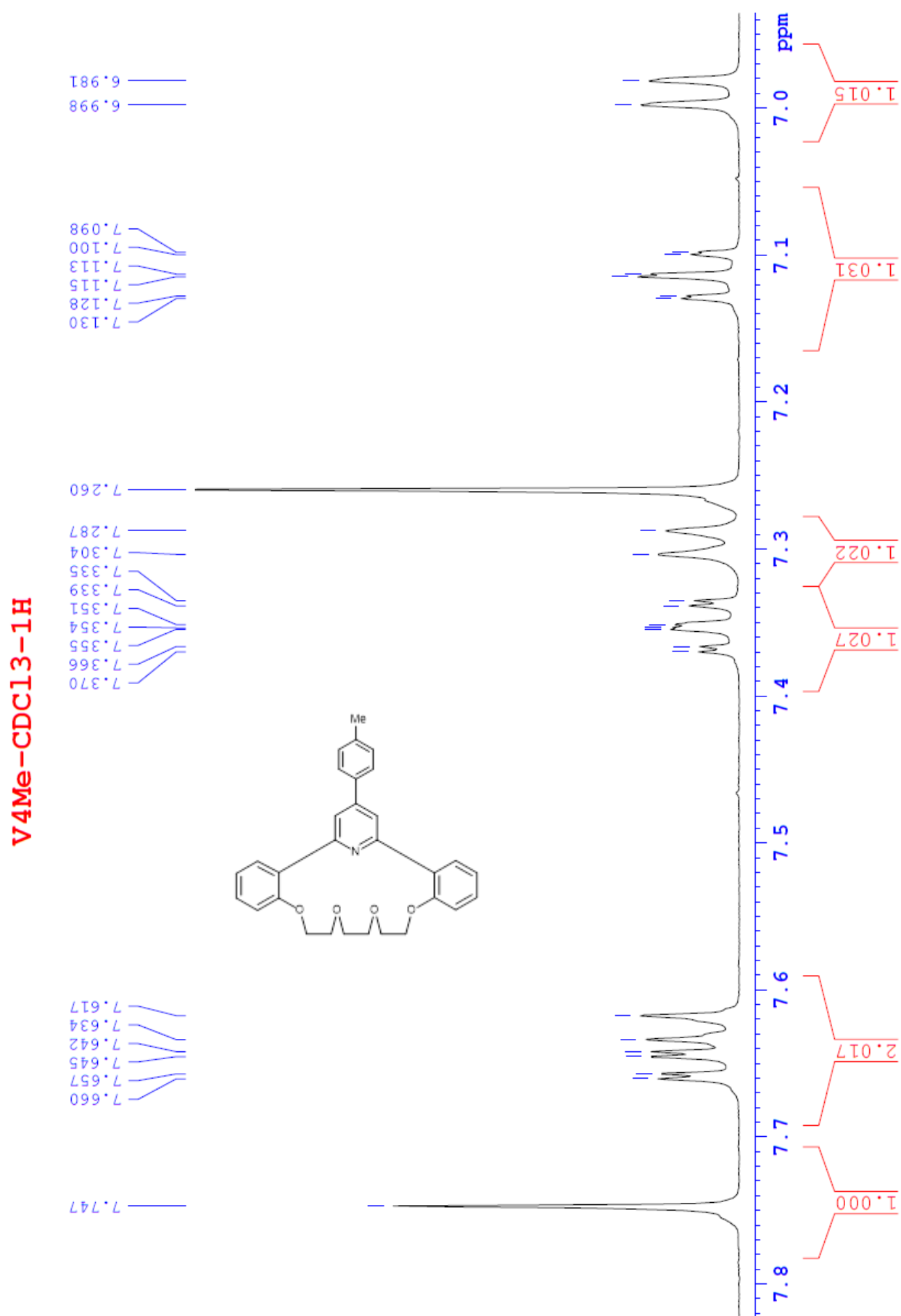
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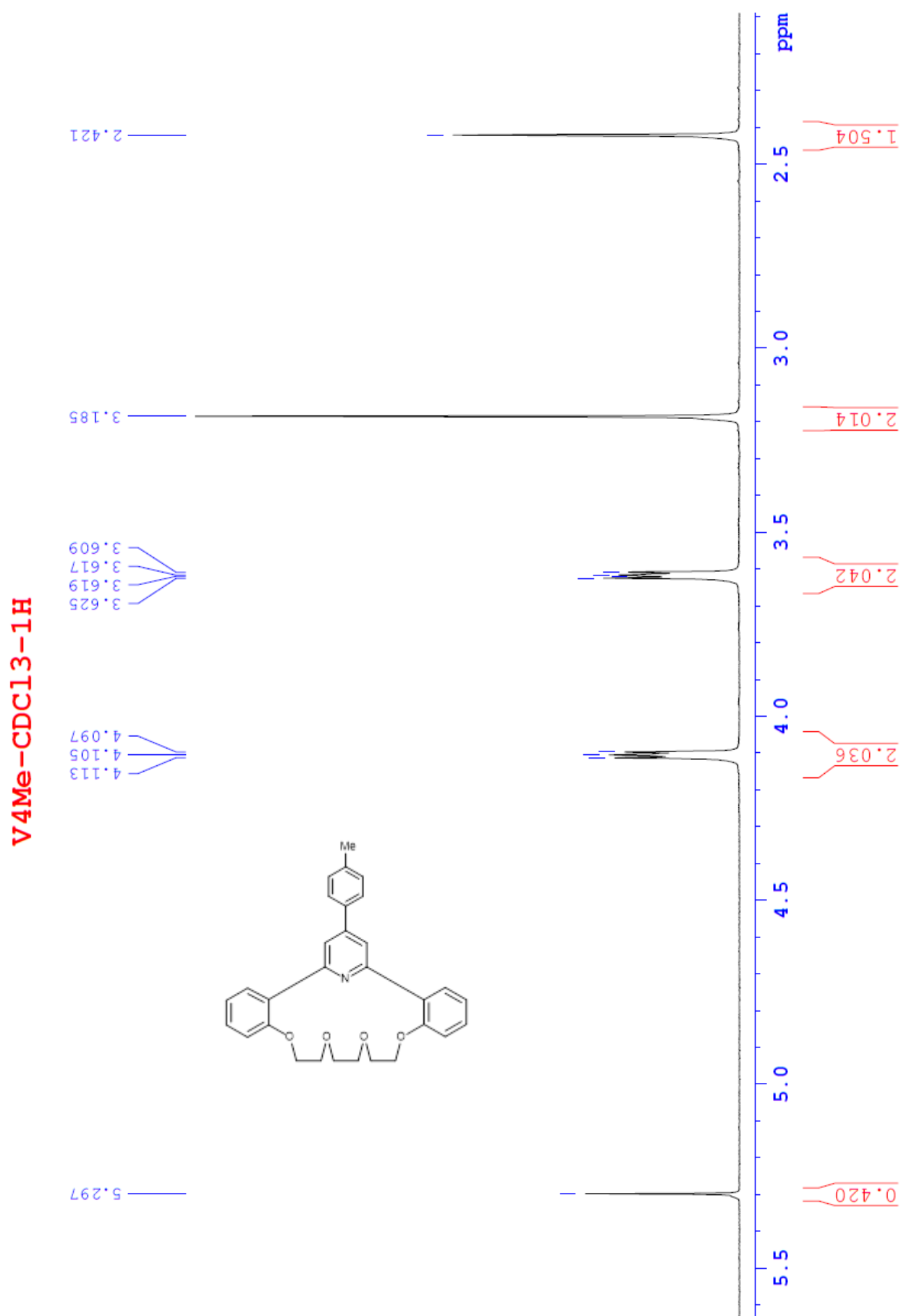
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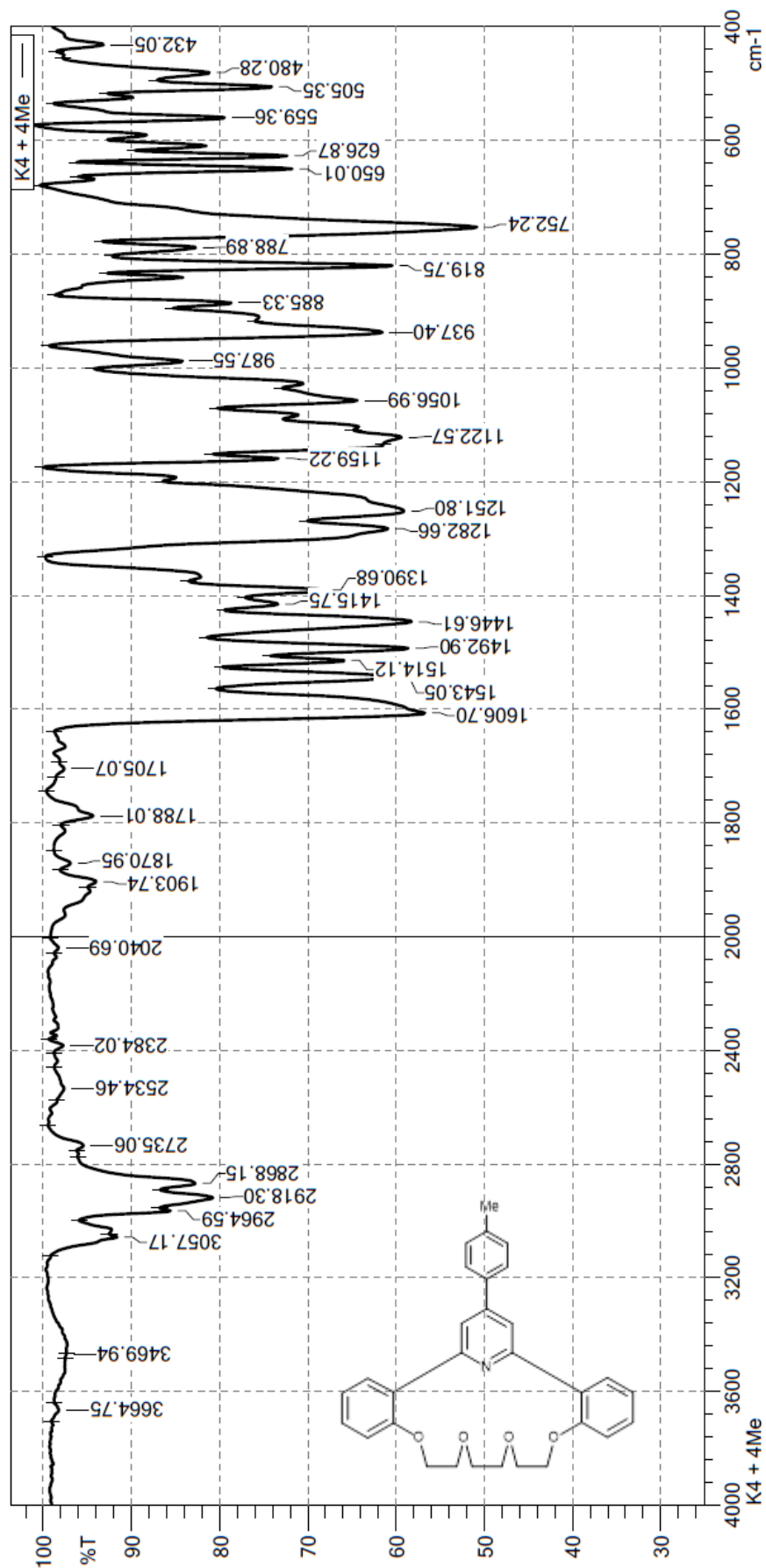




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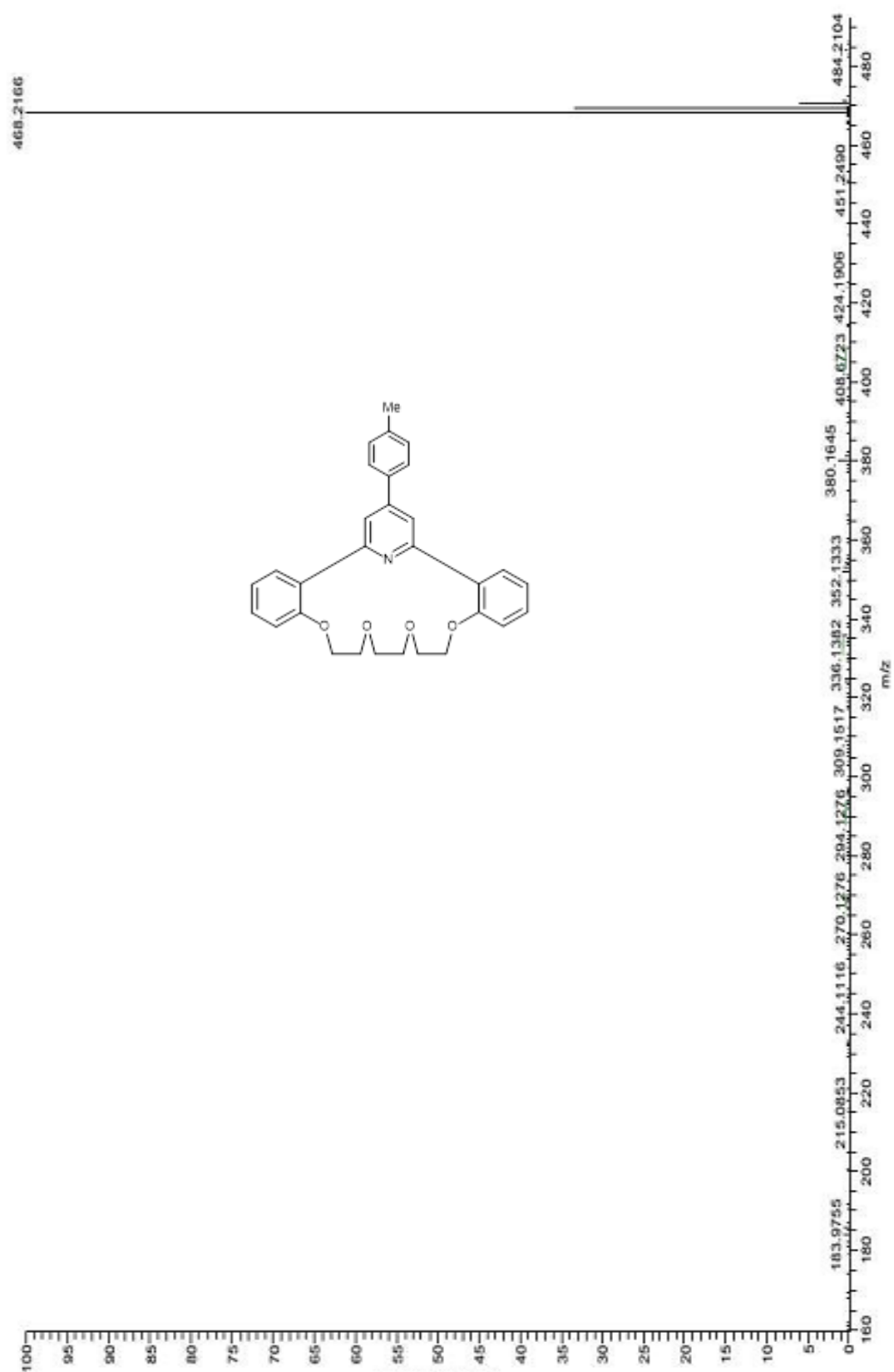
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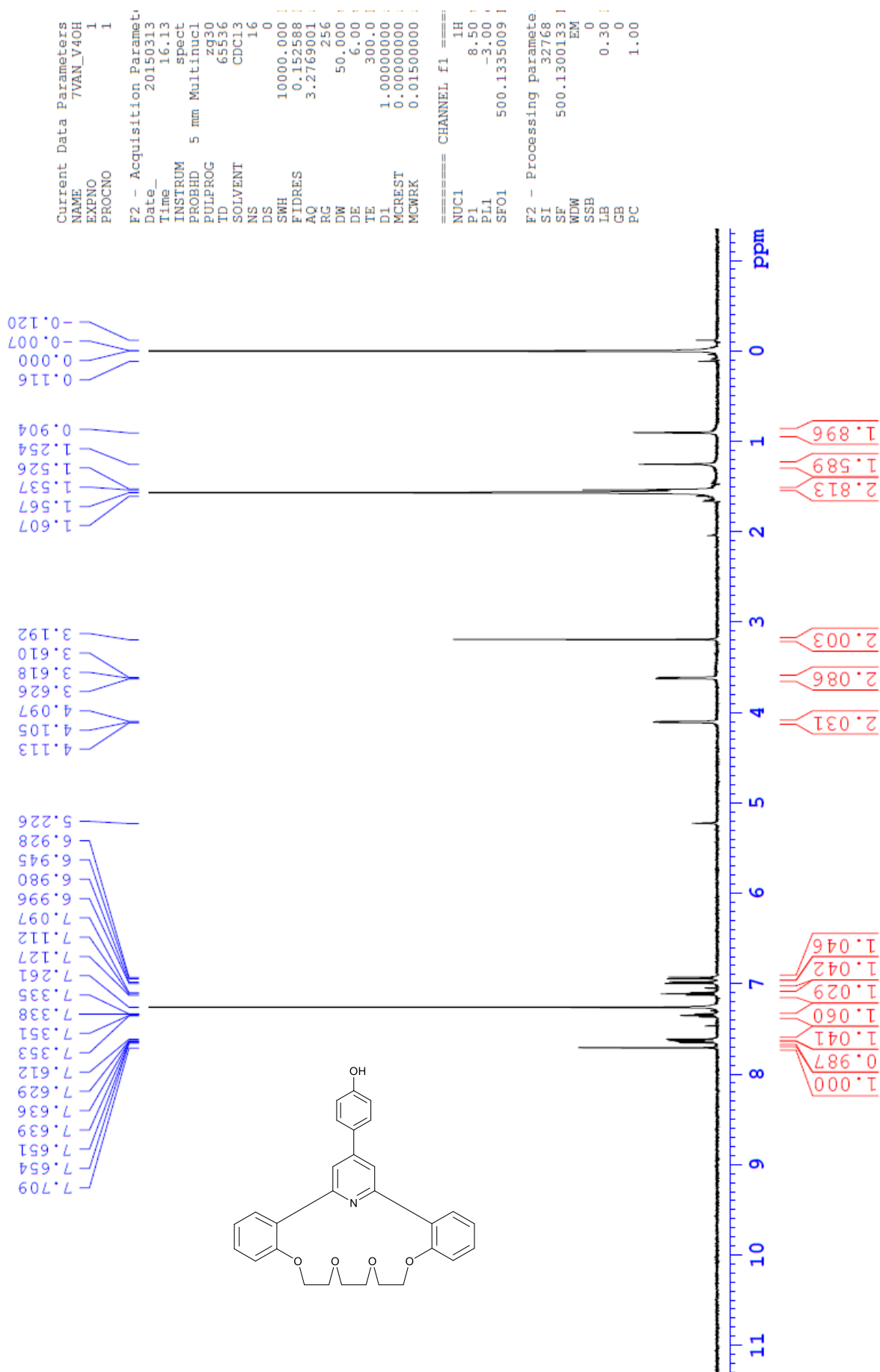
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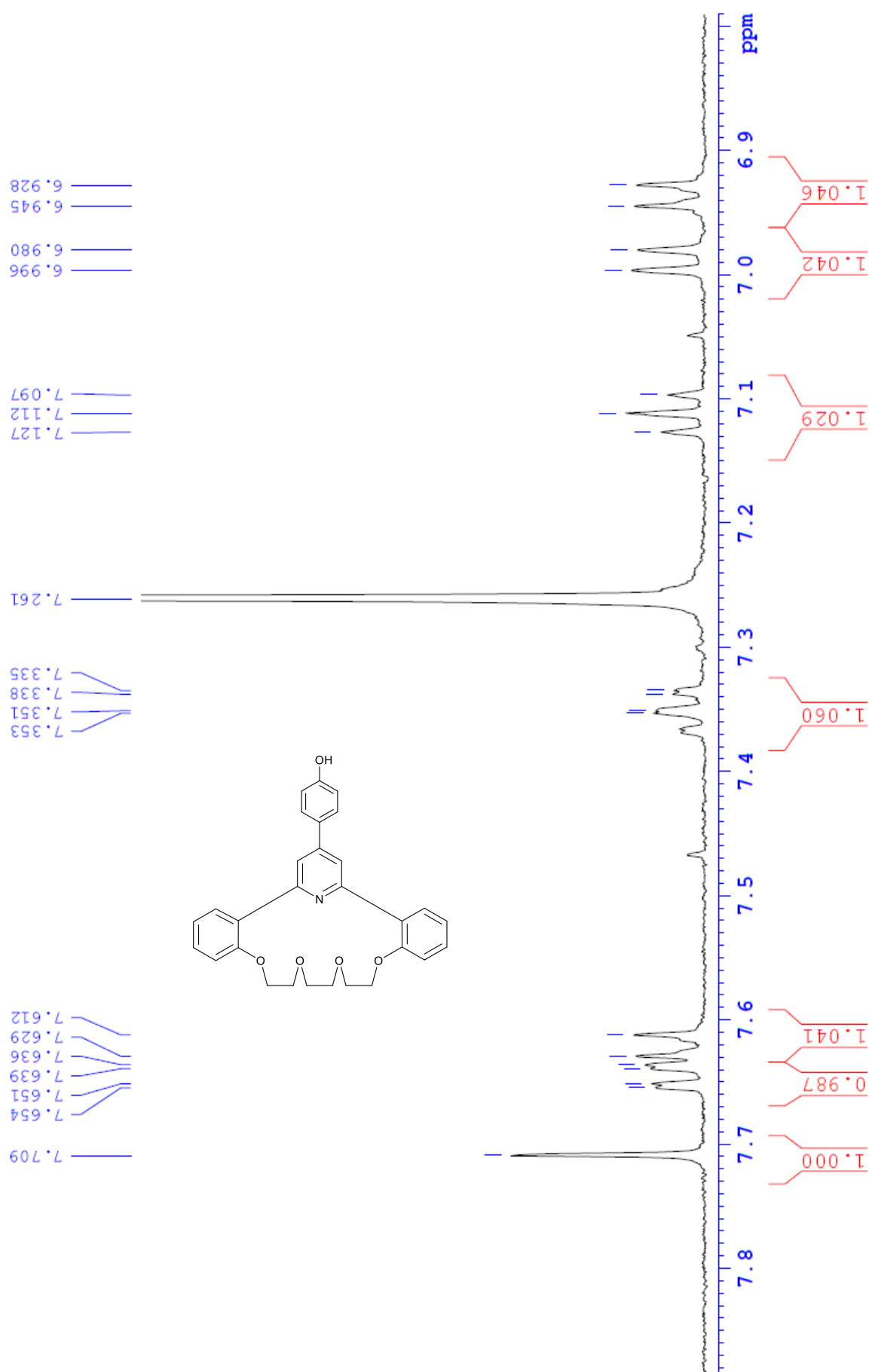
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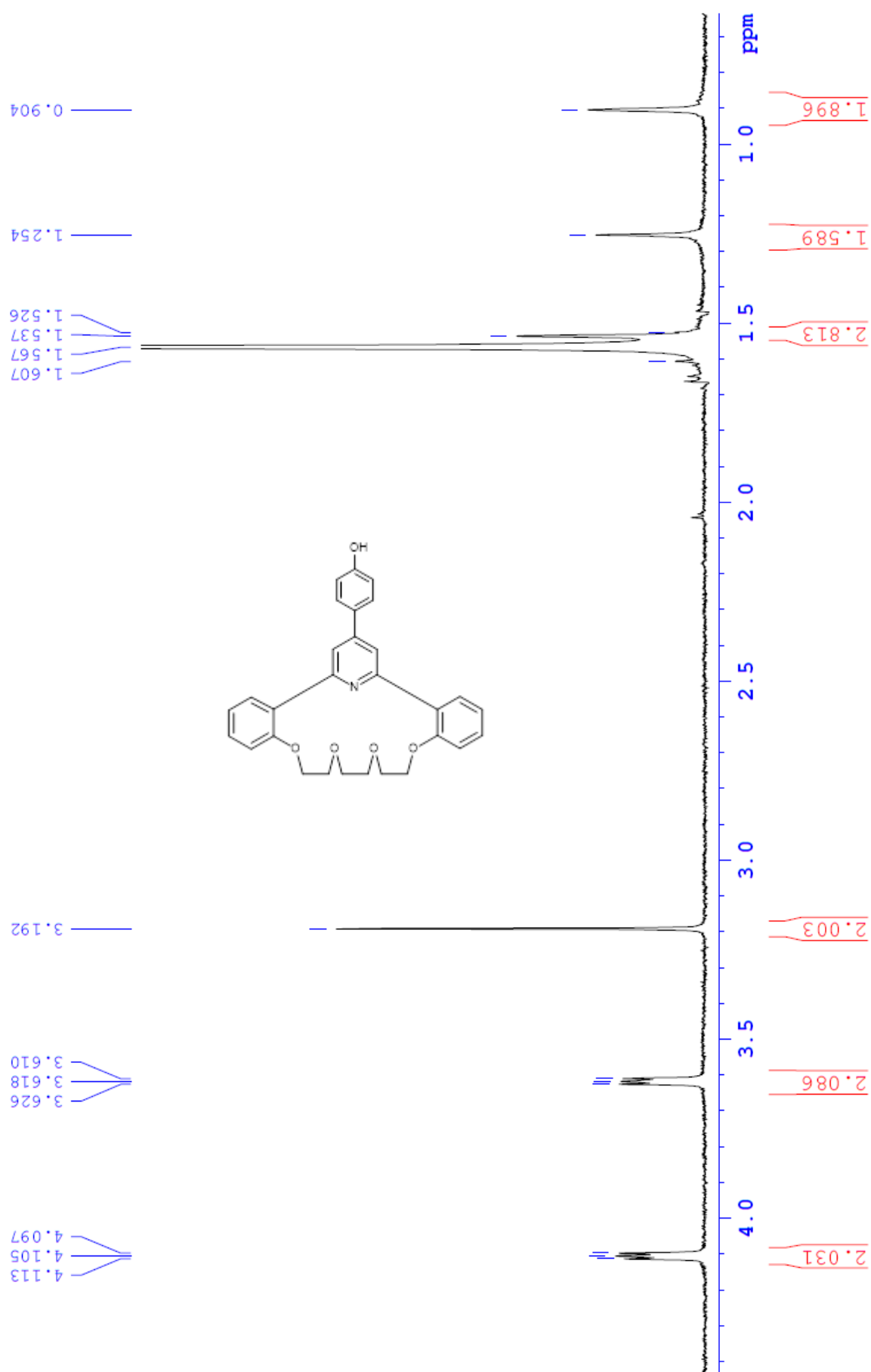
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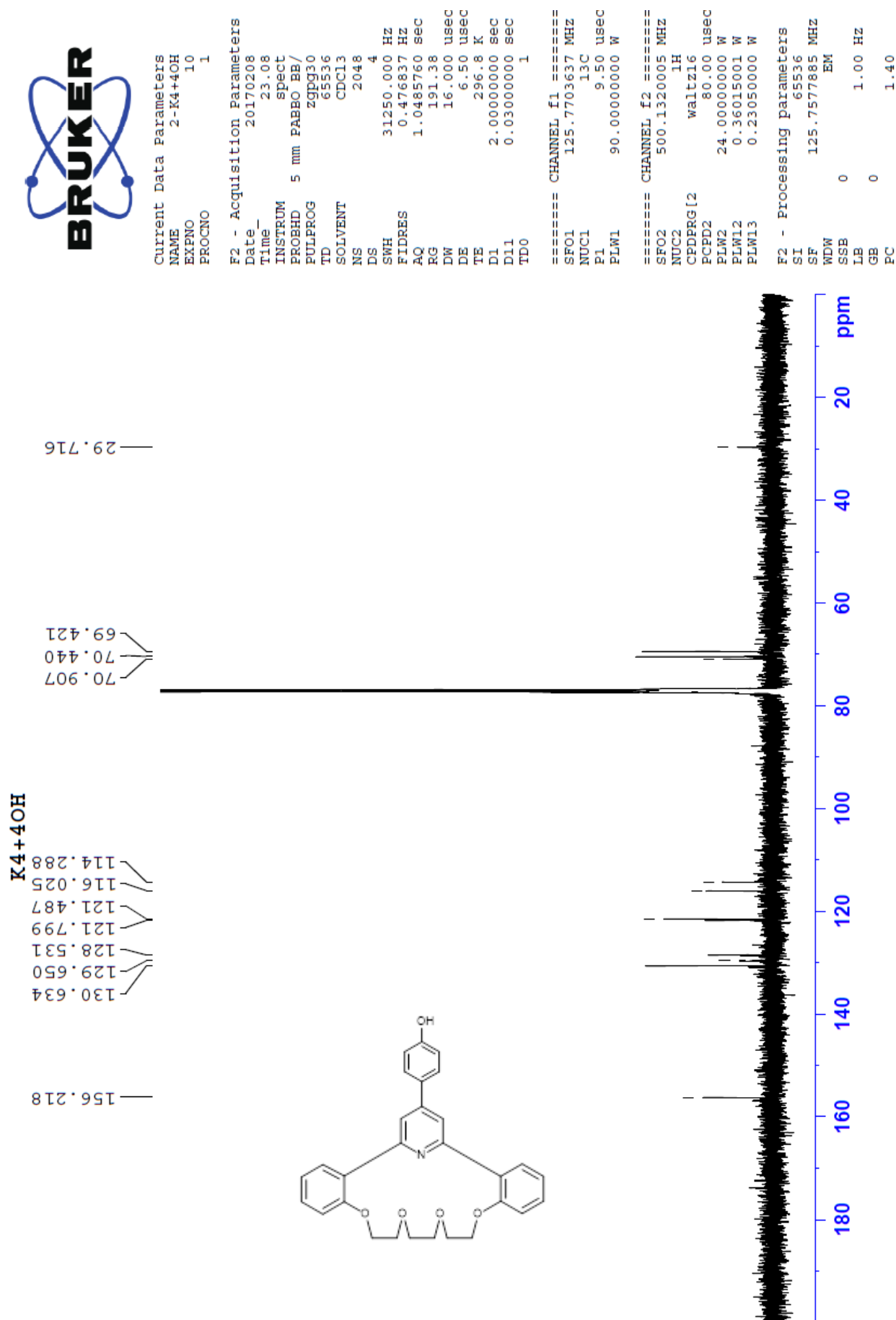
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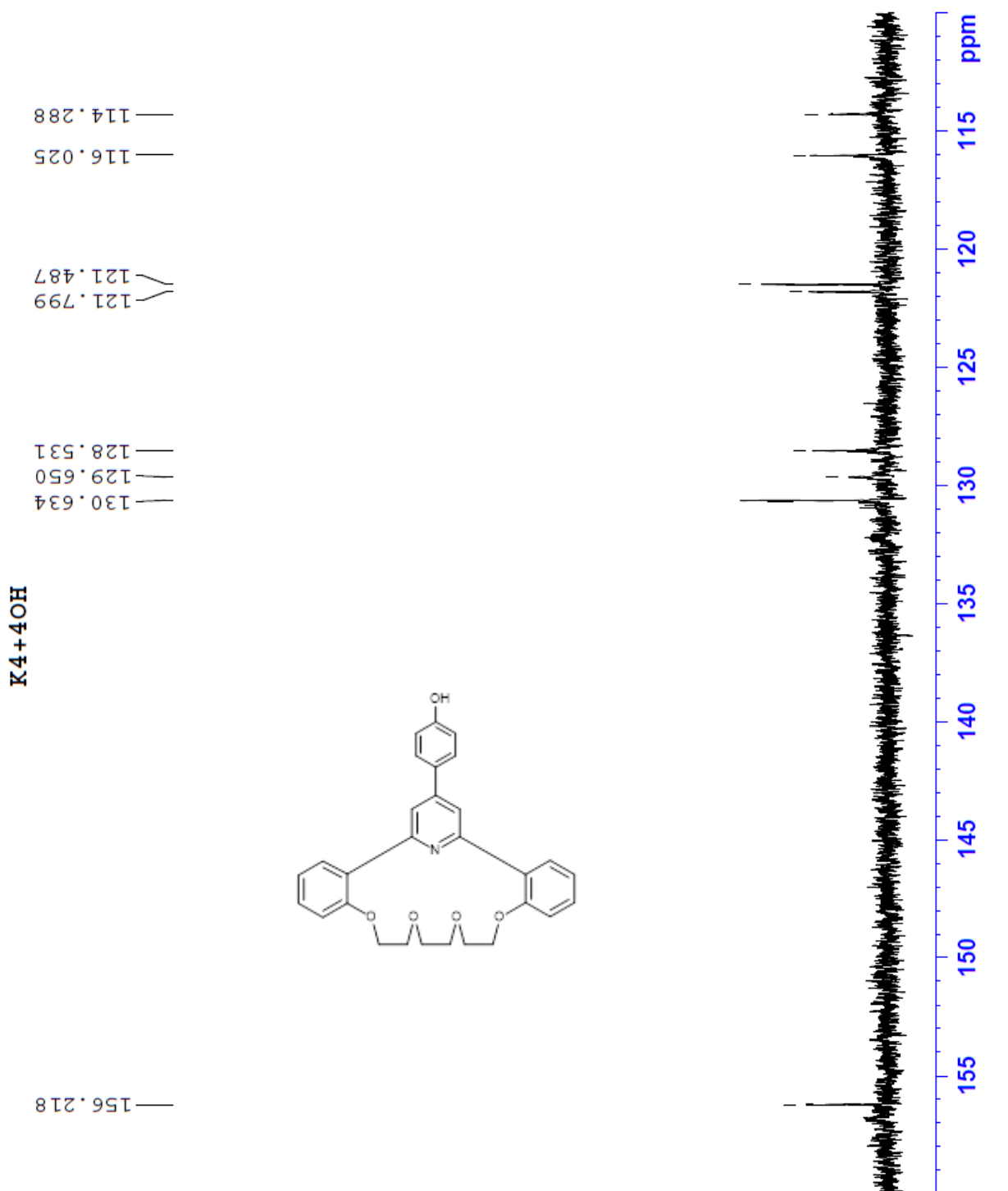


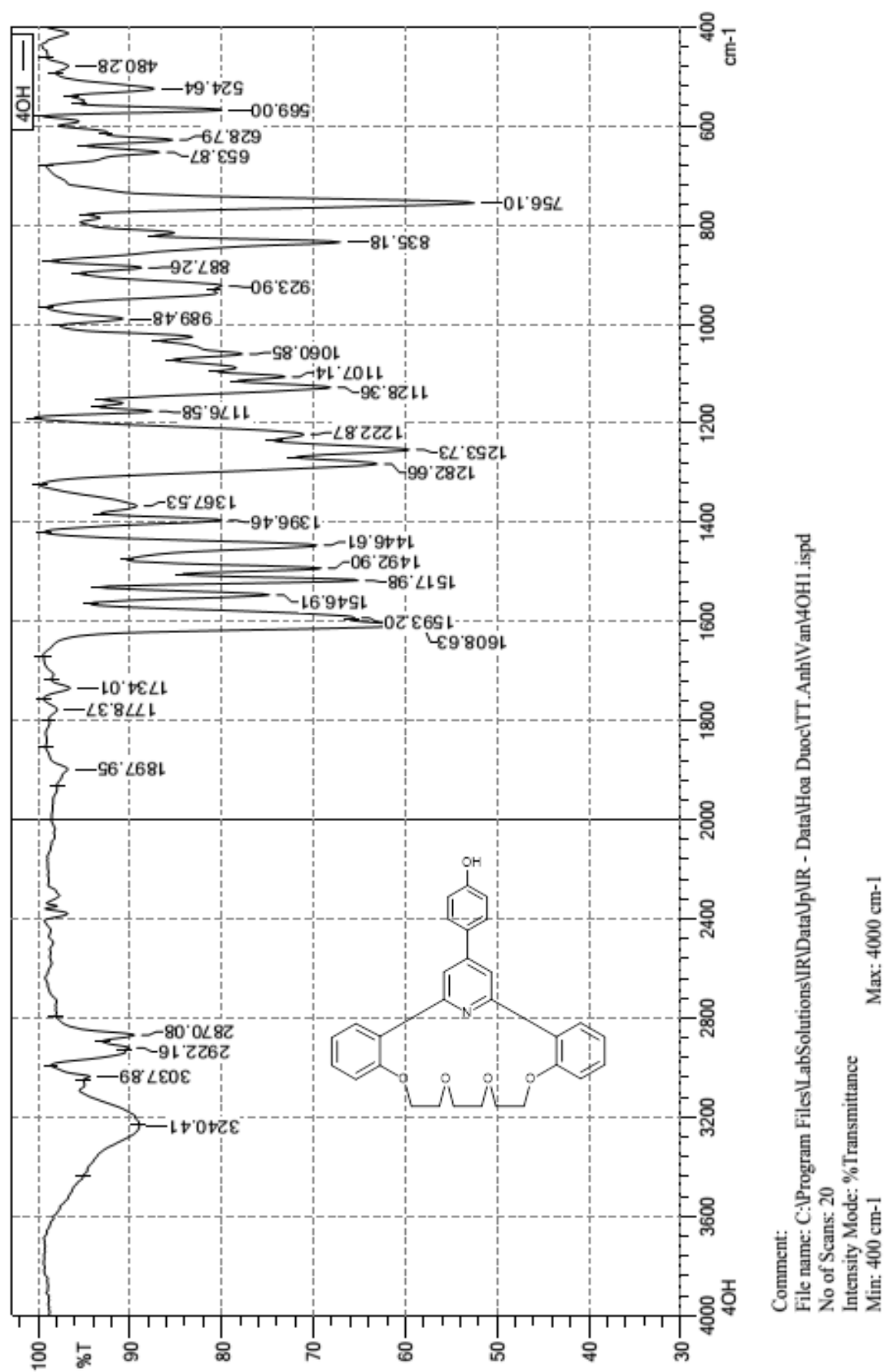
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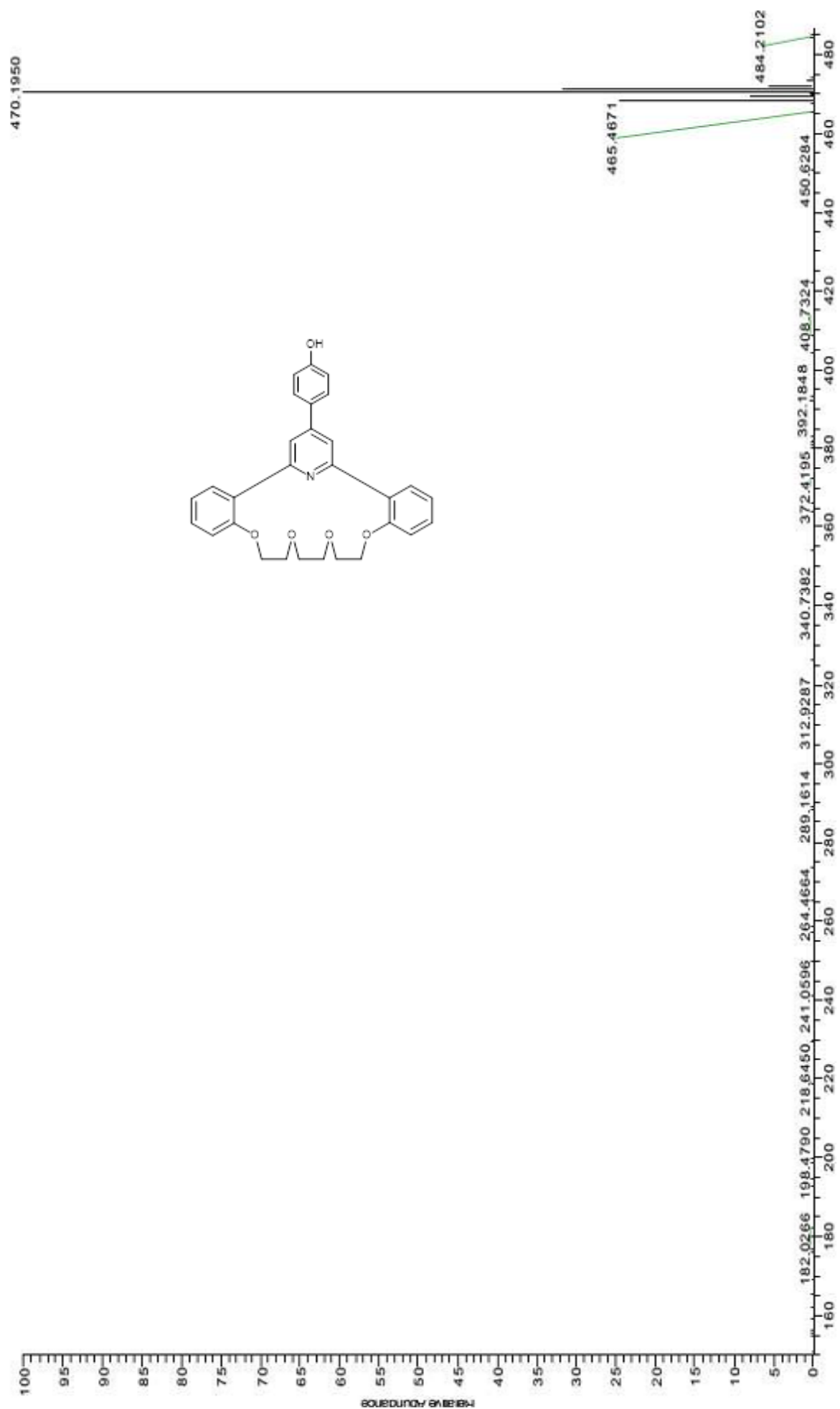
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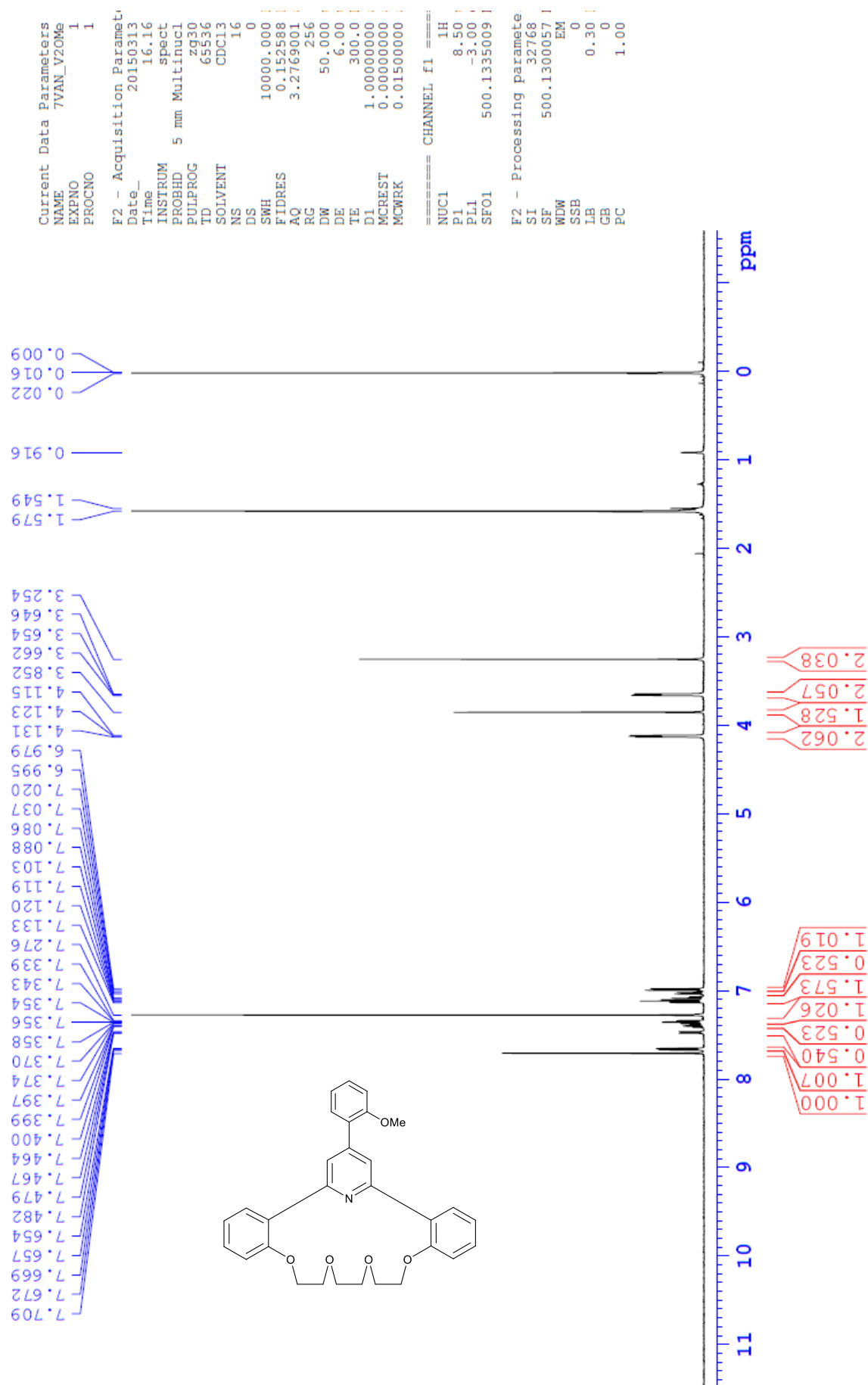


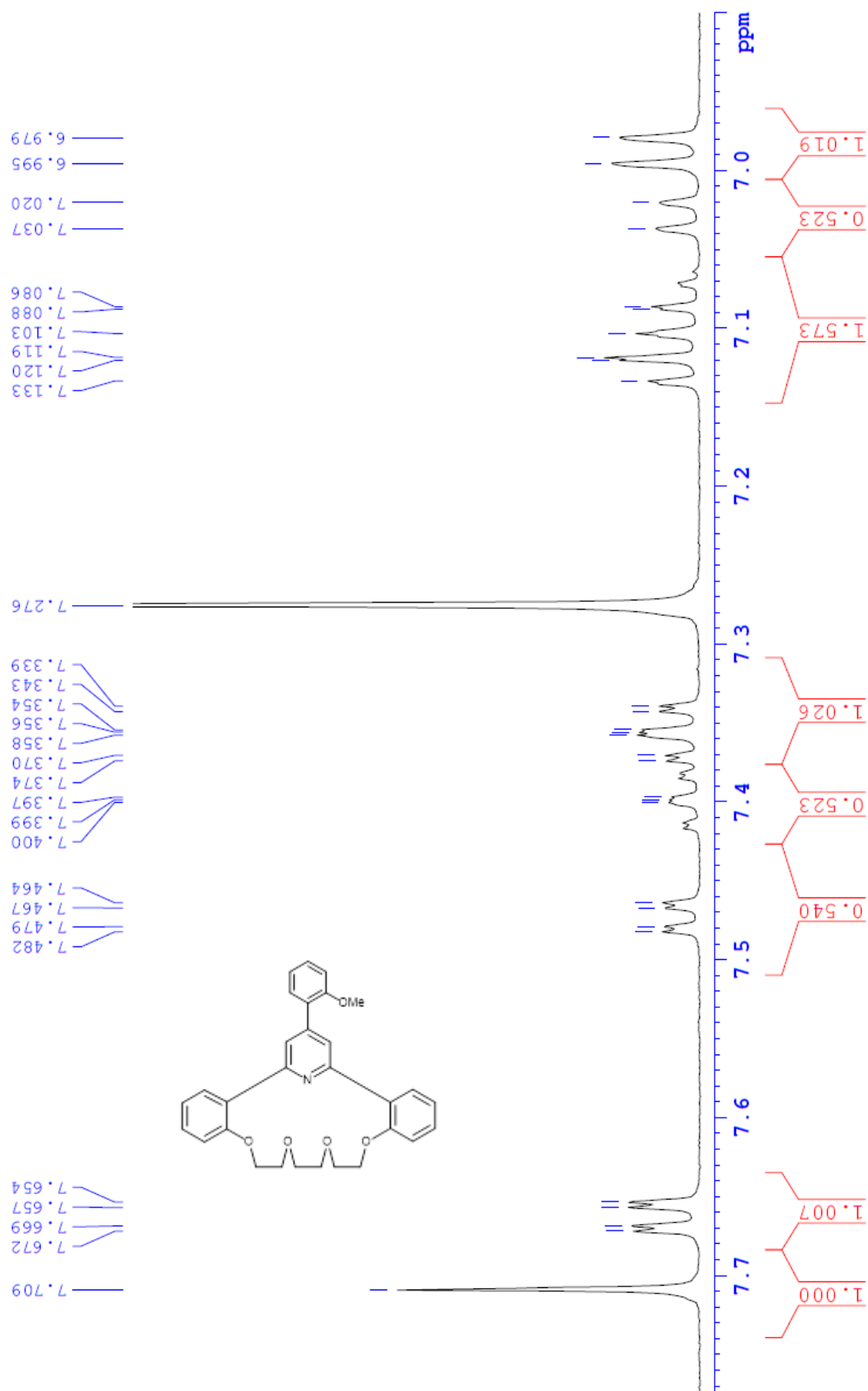


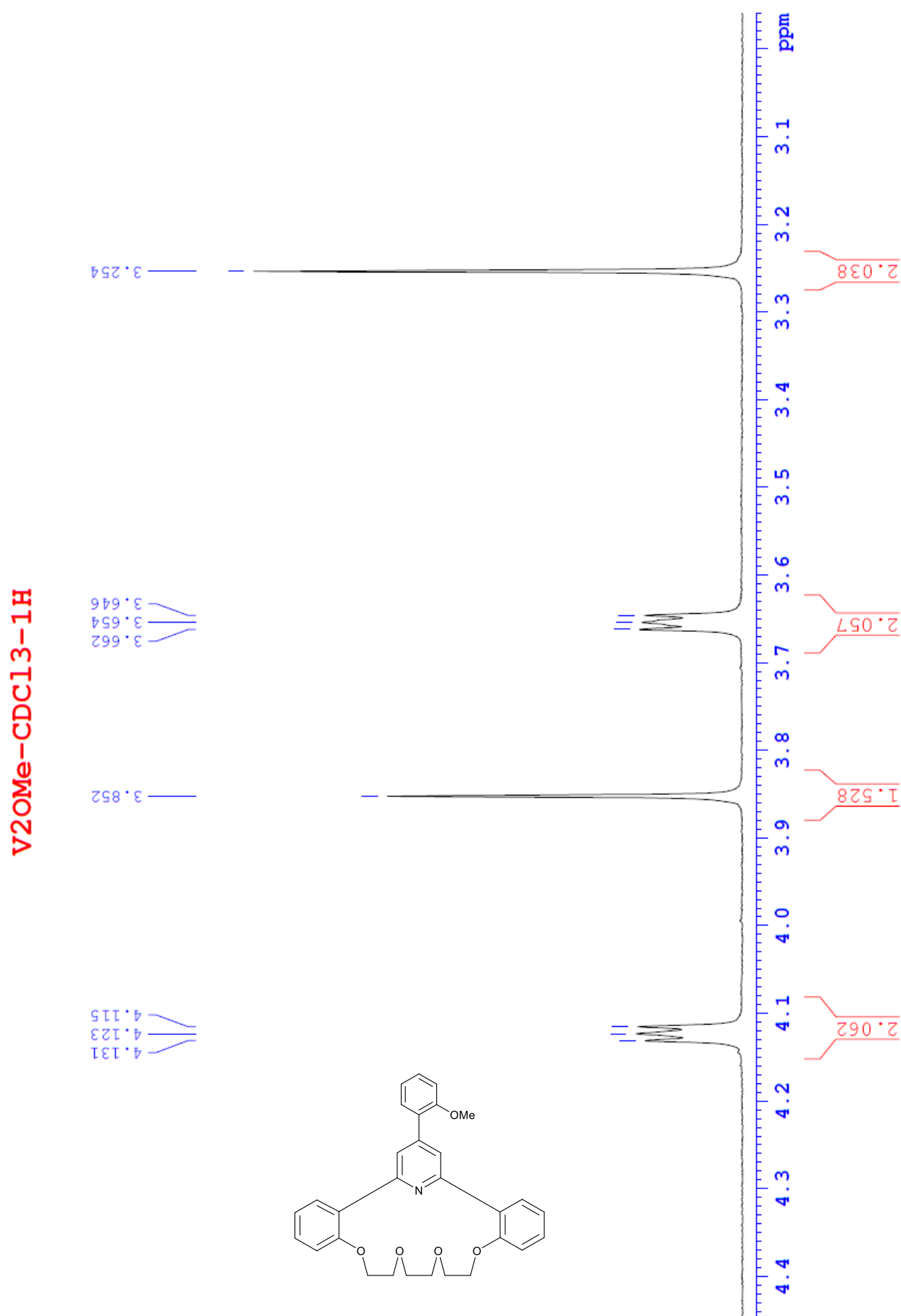




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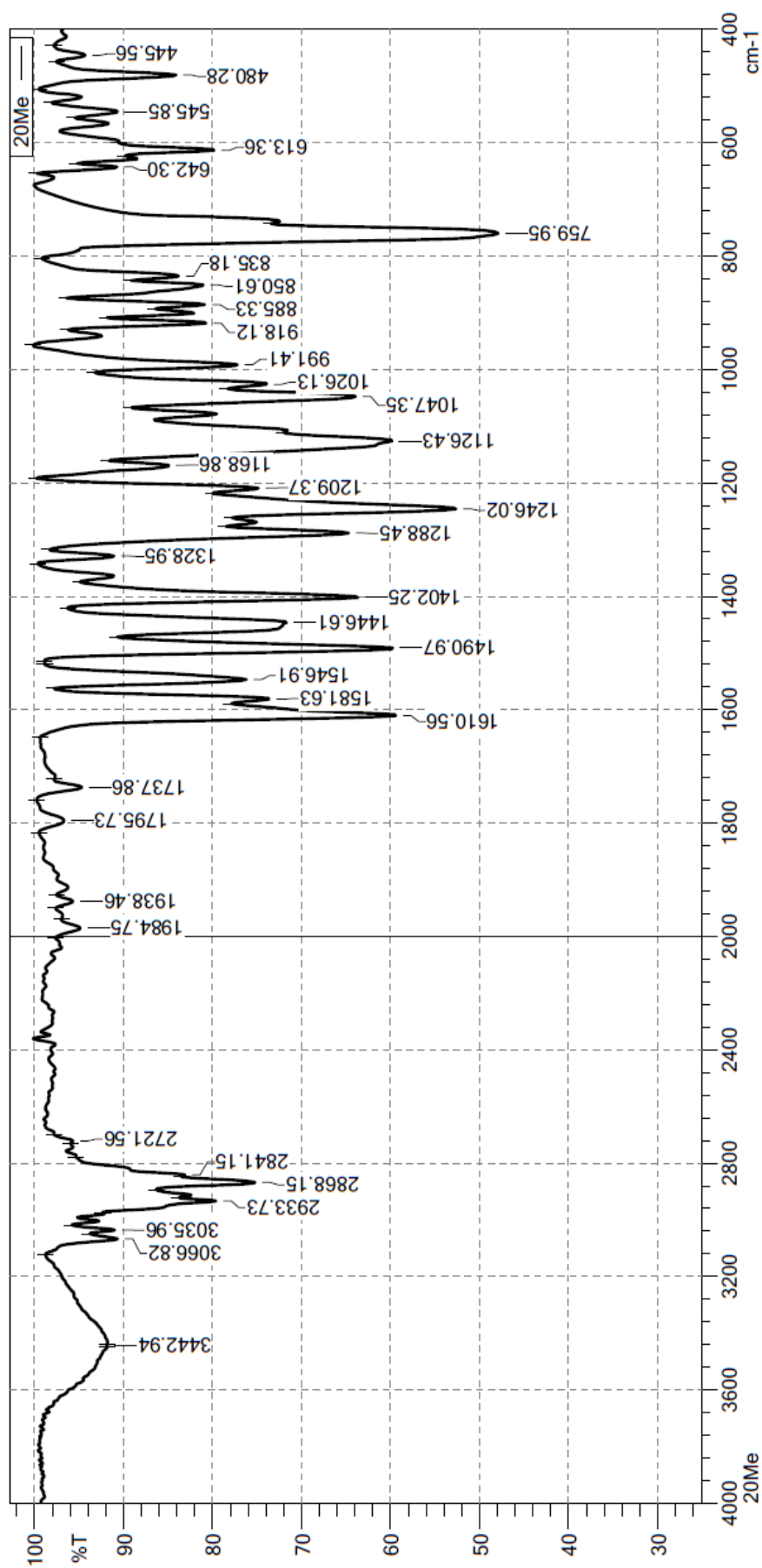


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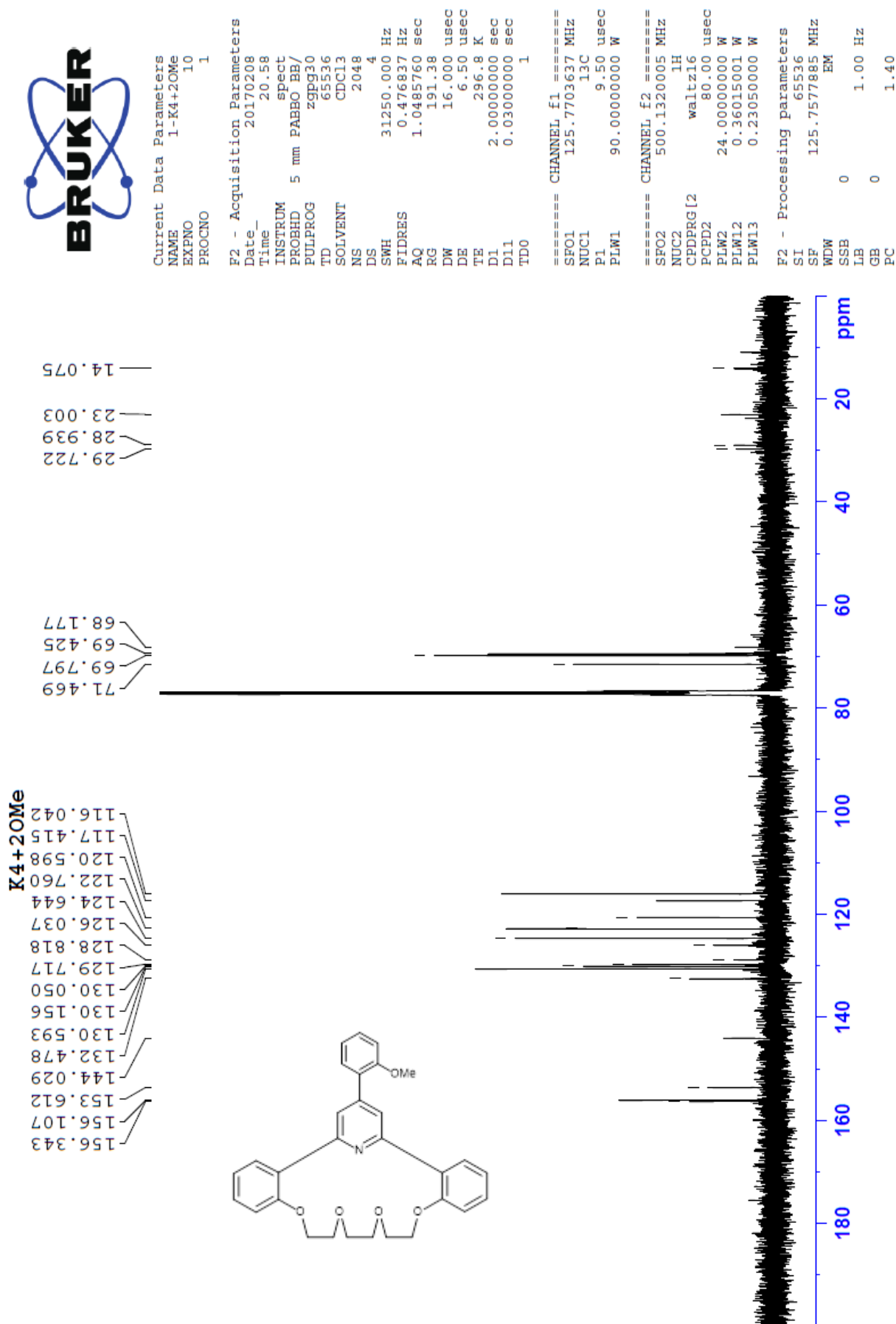


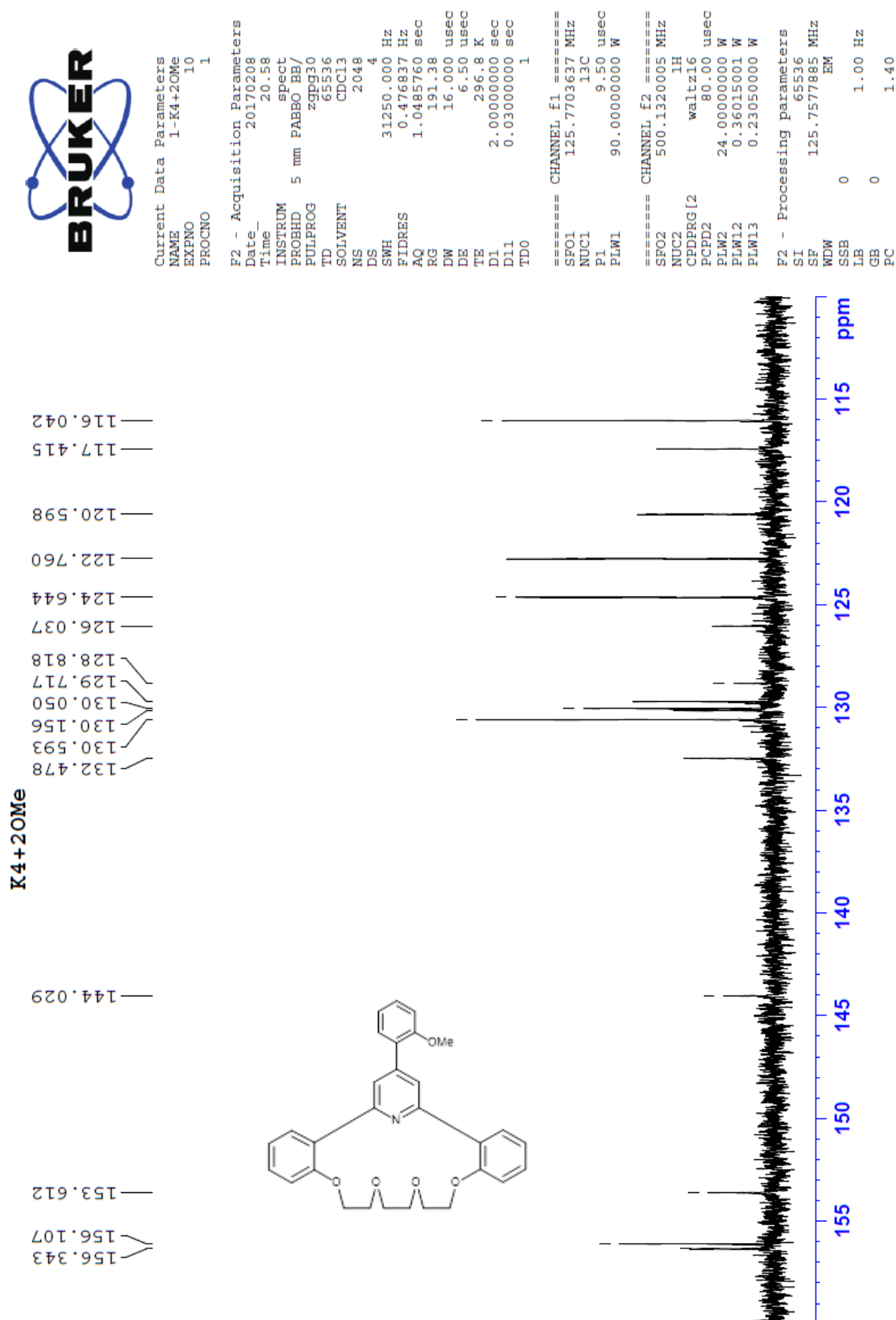
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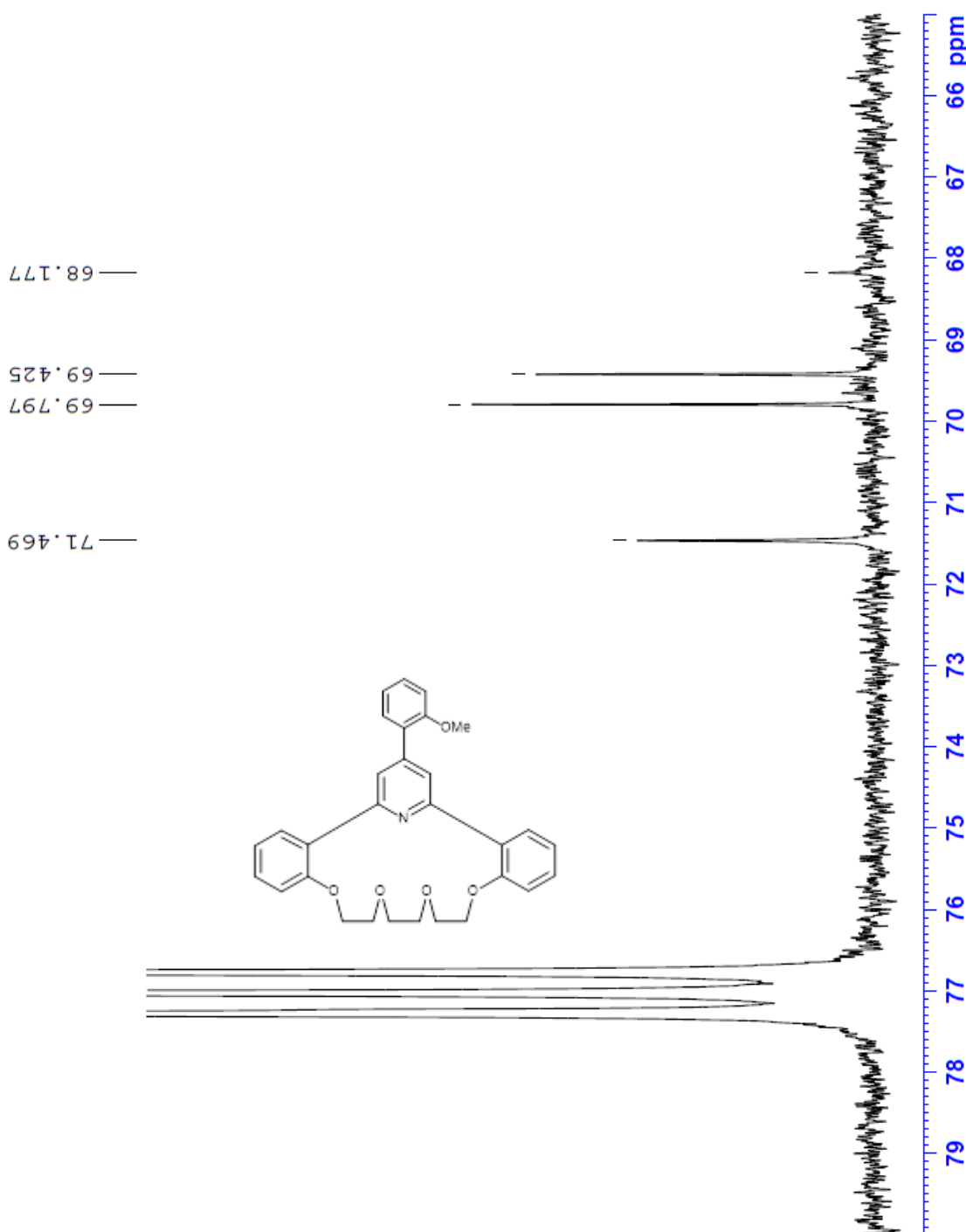
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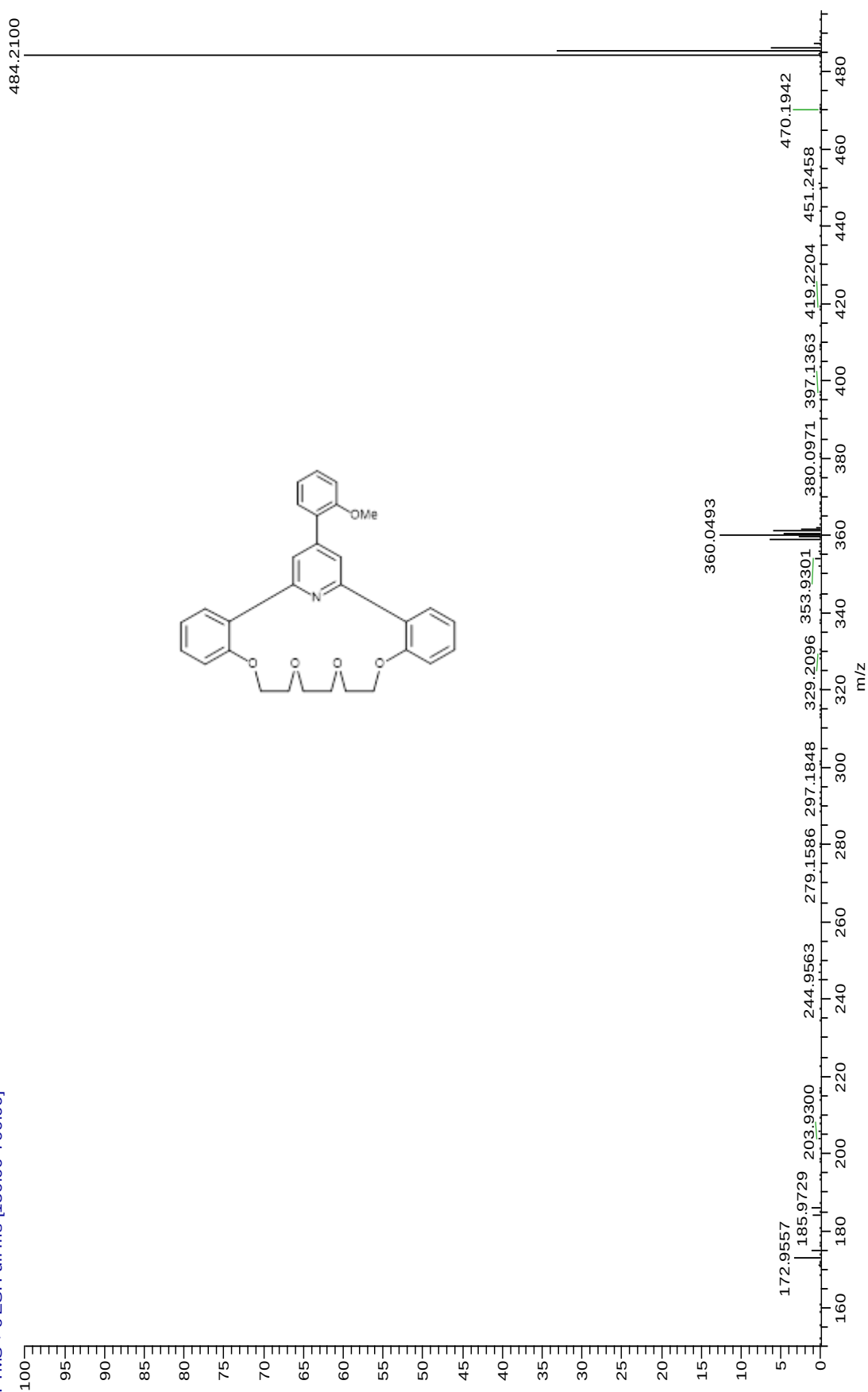
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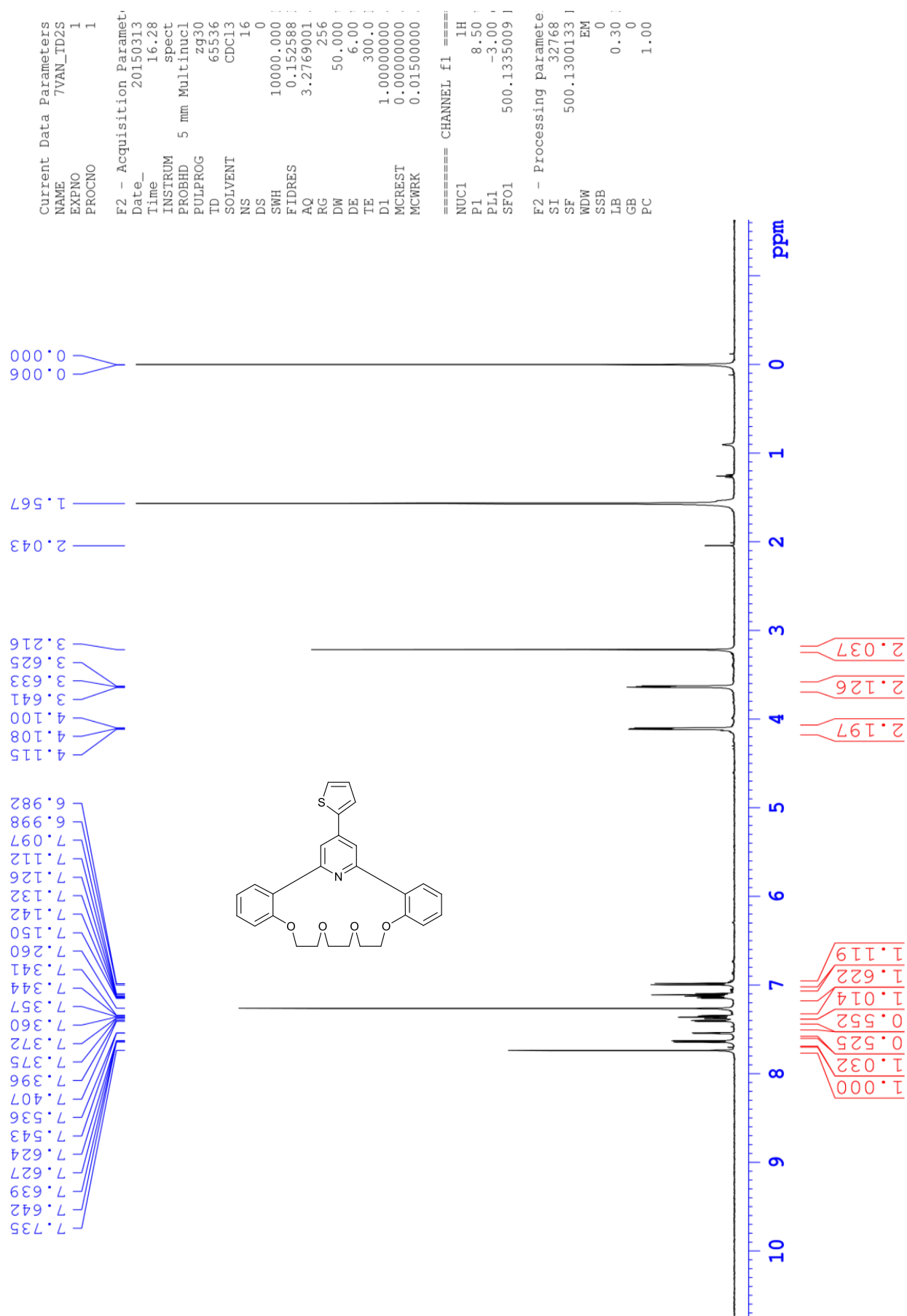
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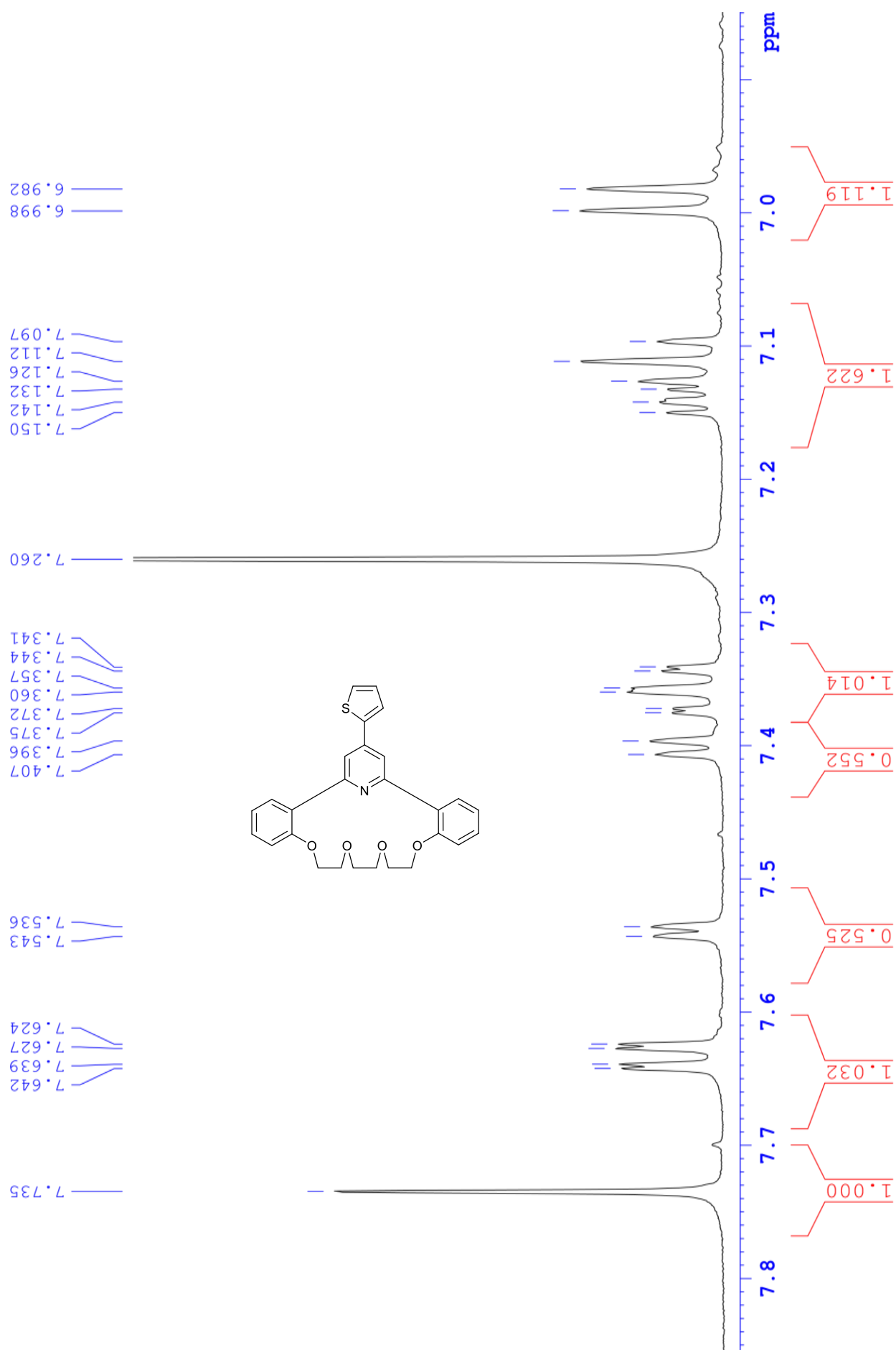
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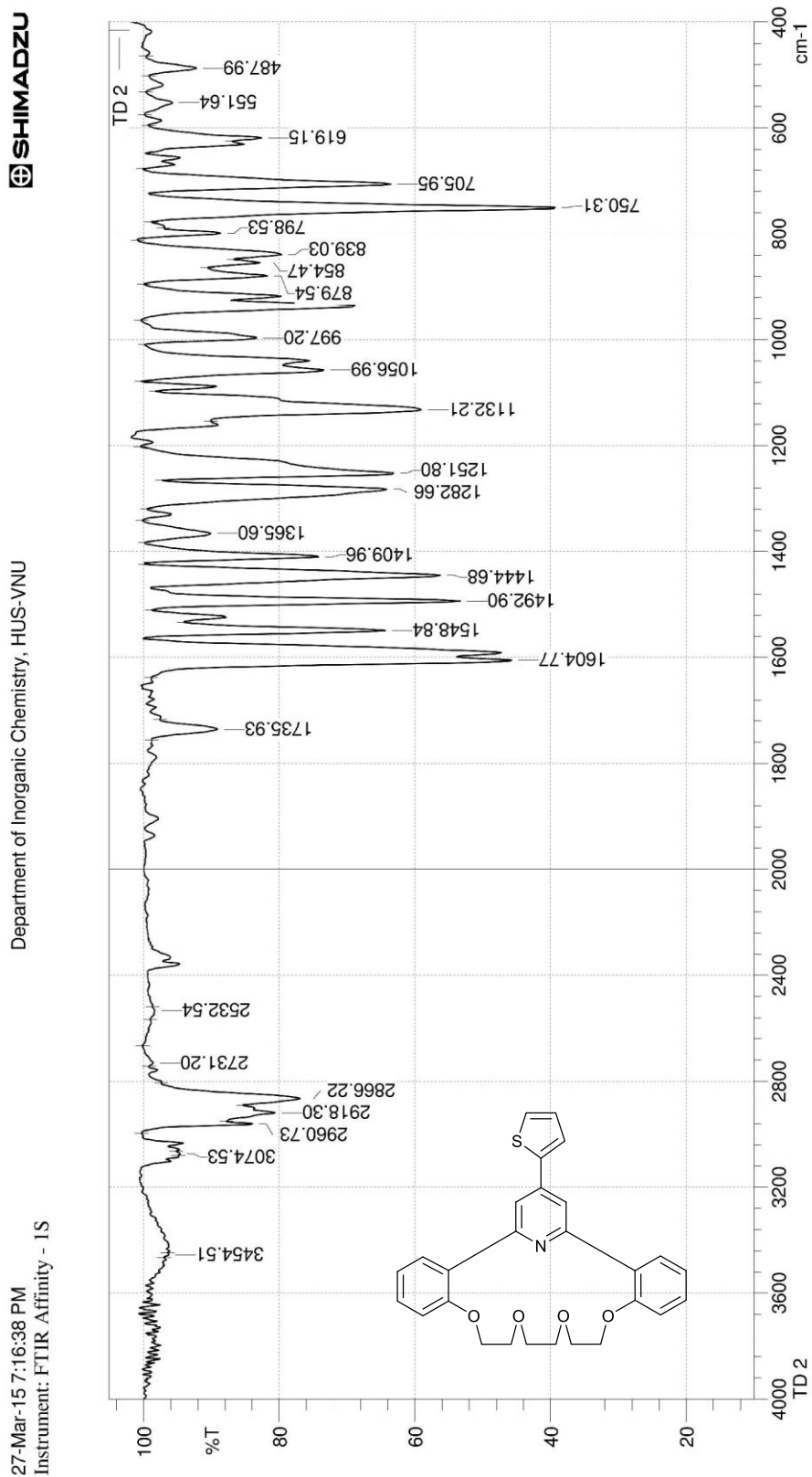


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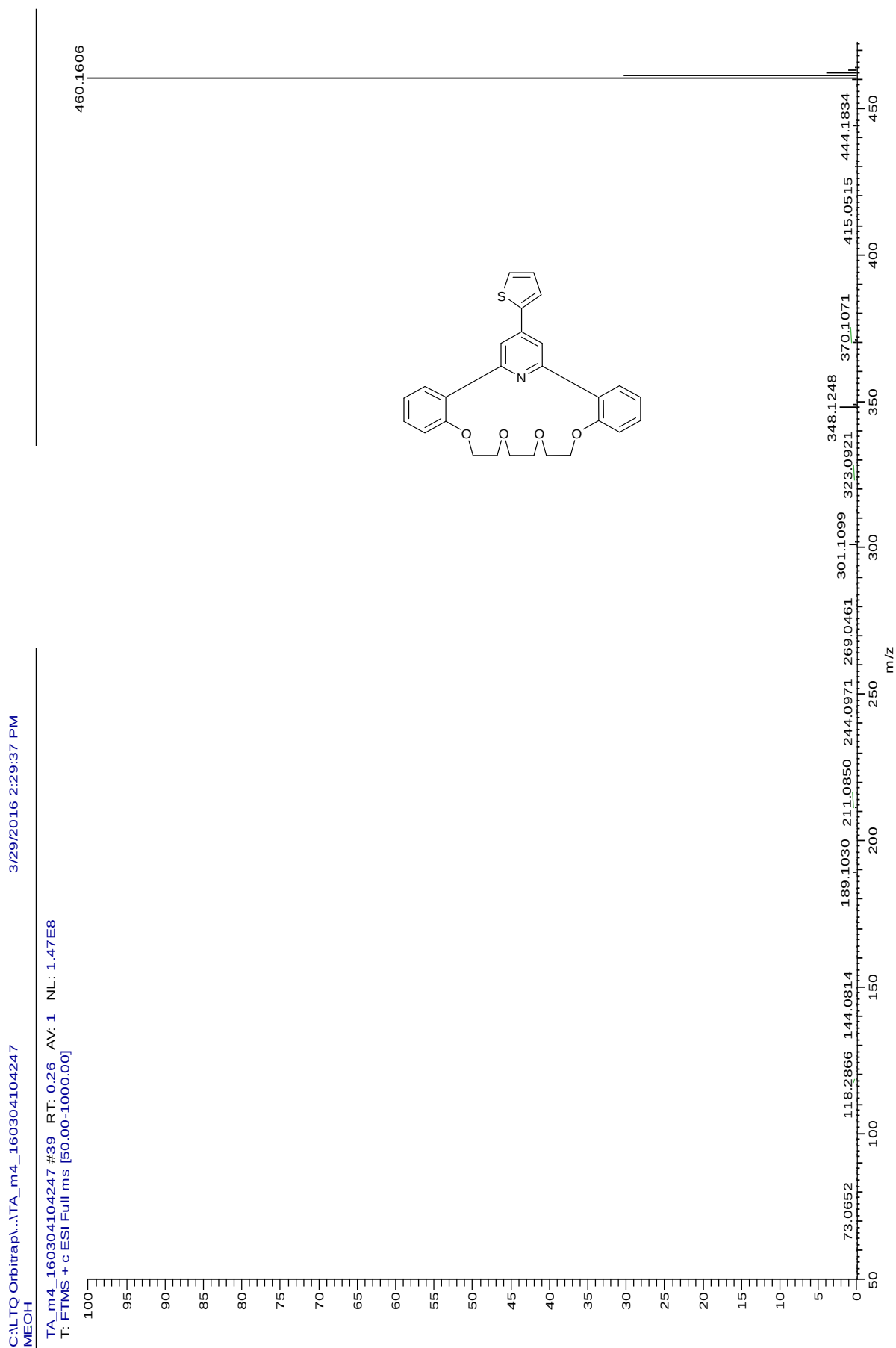
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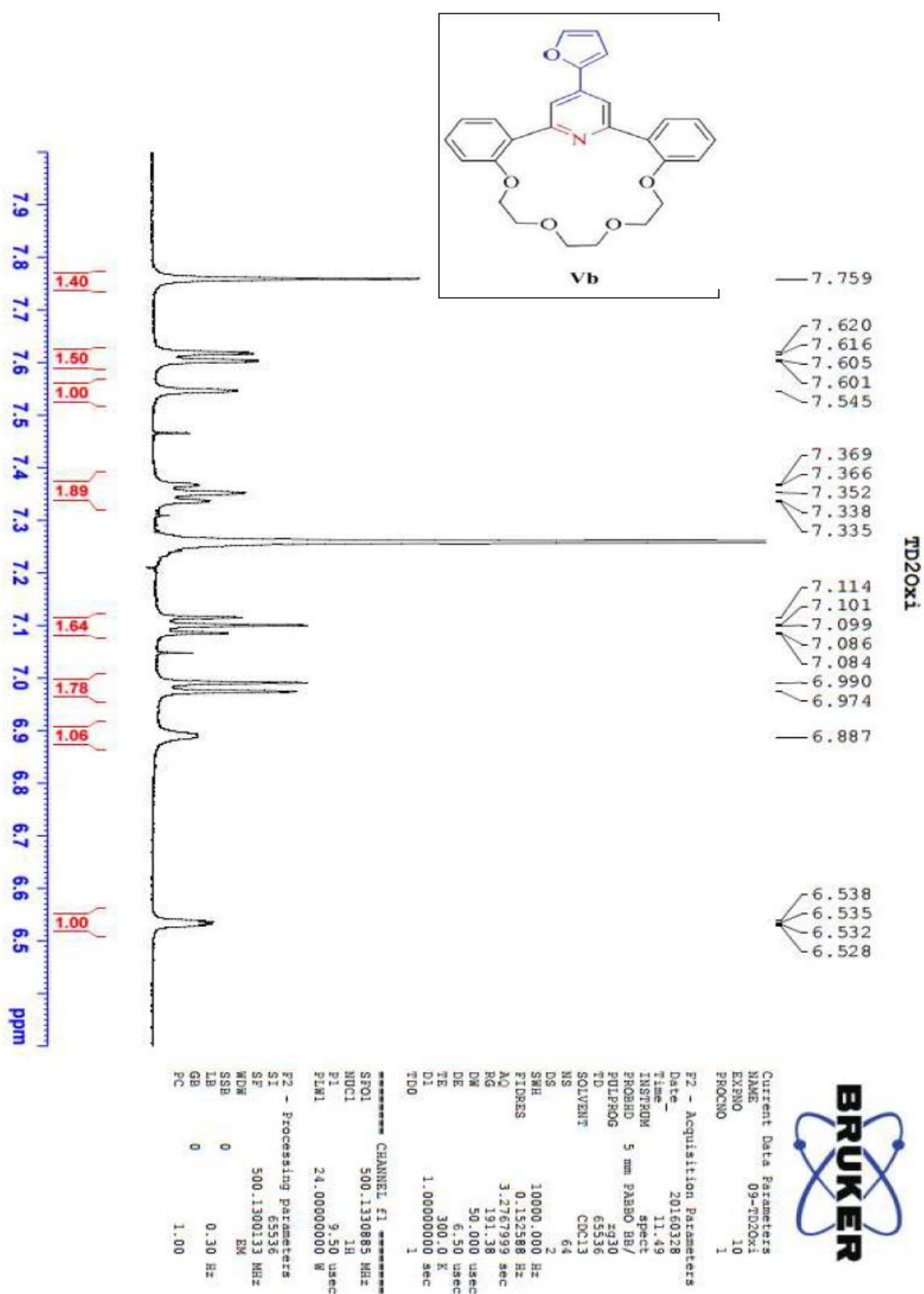


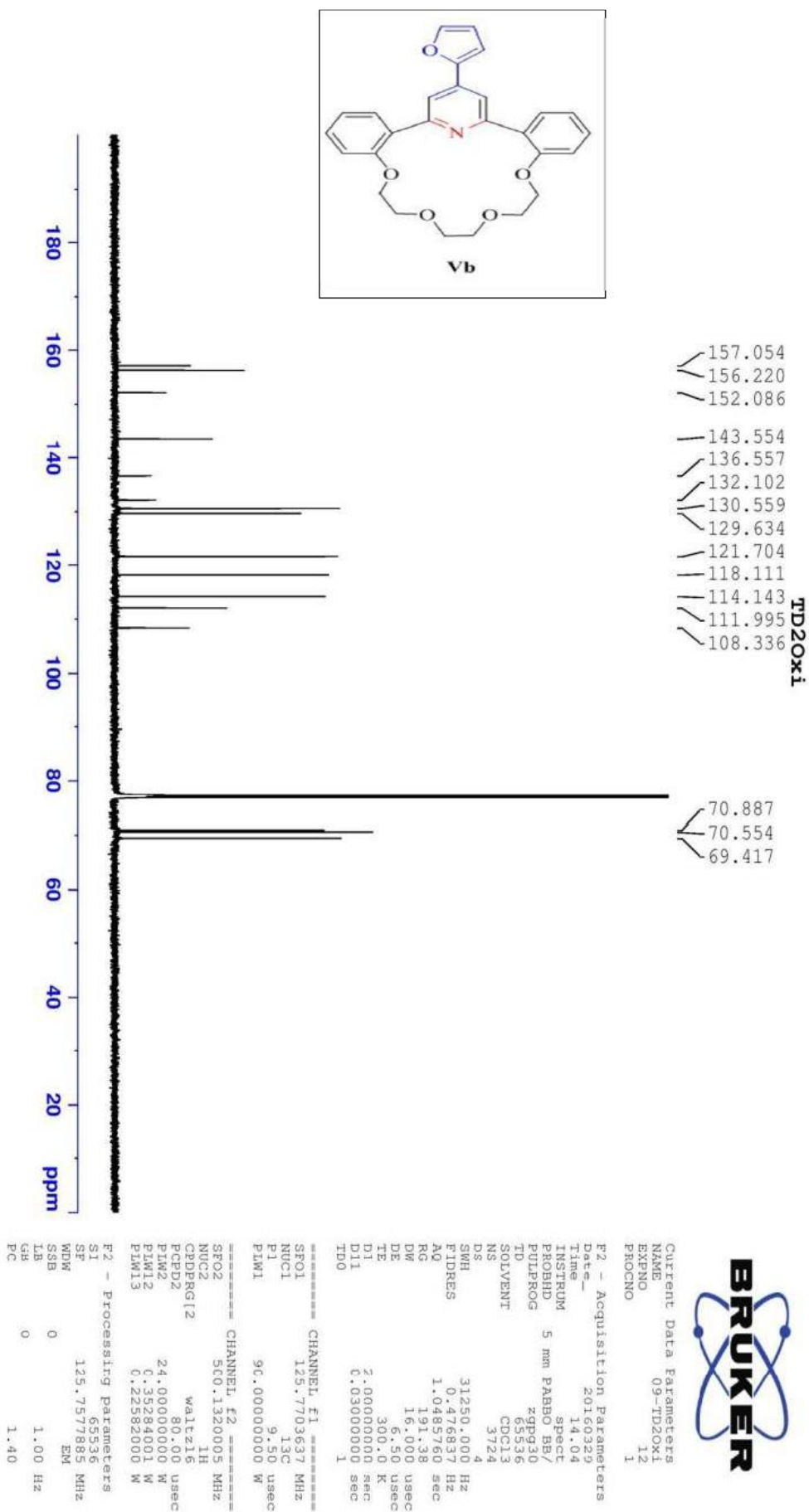


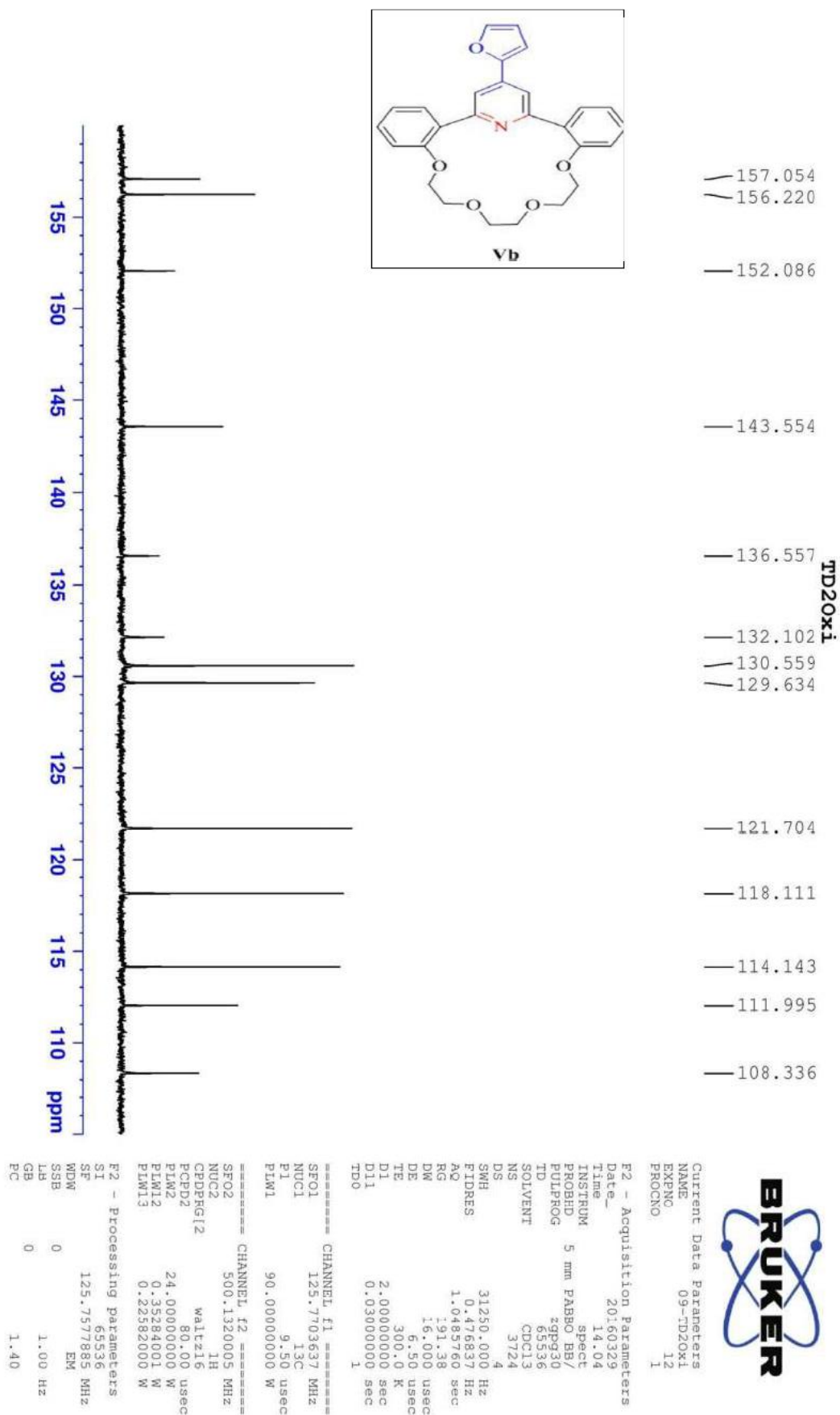


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Atmosphere Correction:



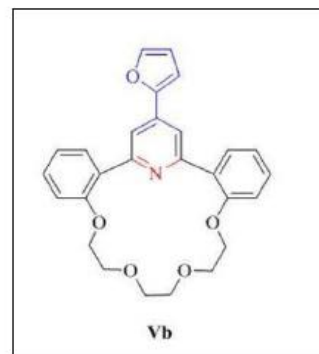
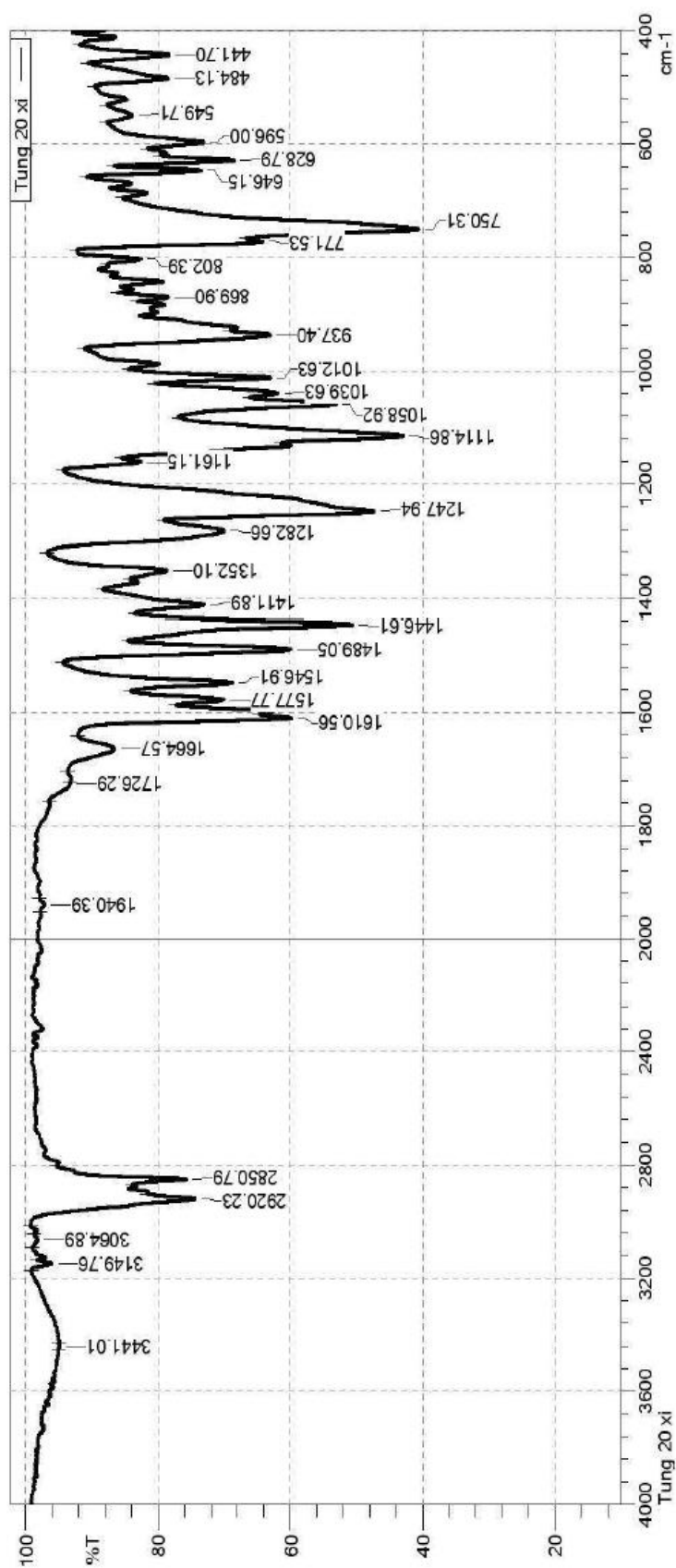






SHIMADZU

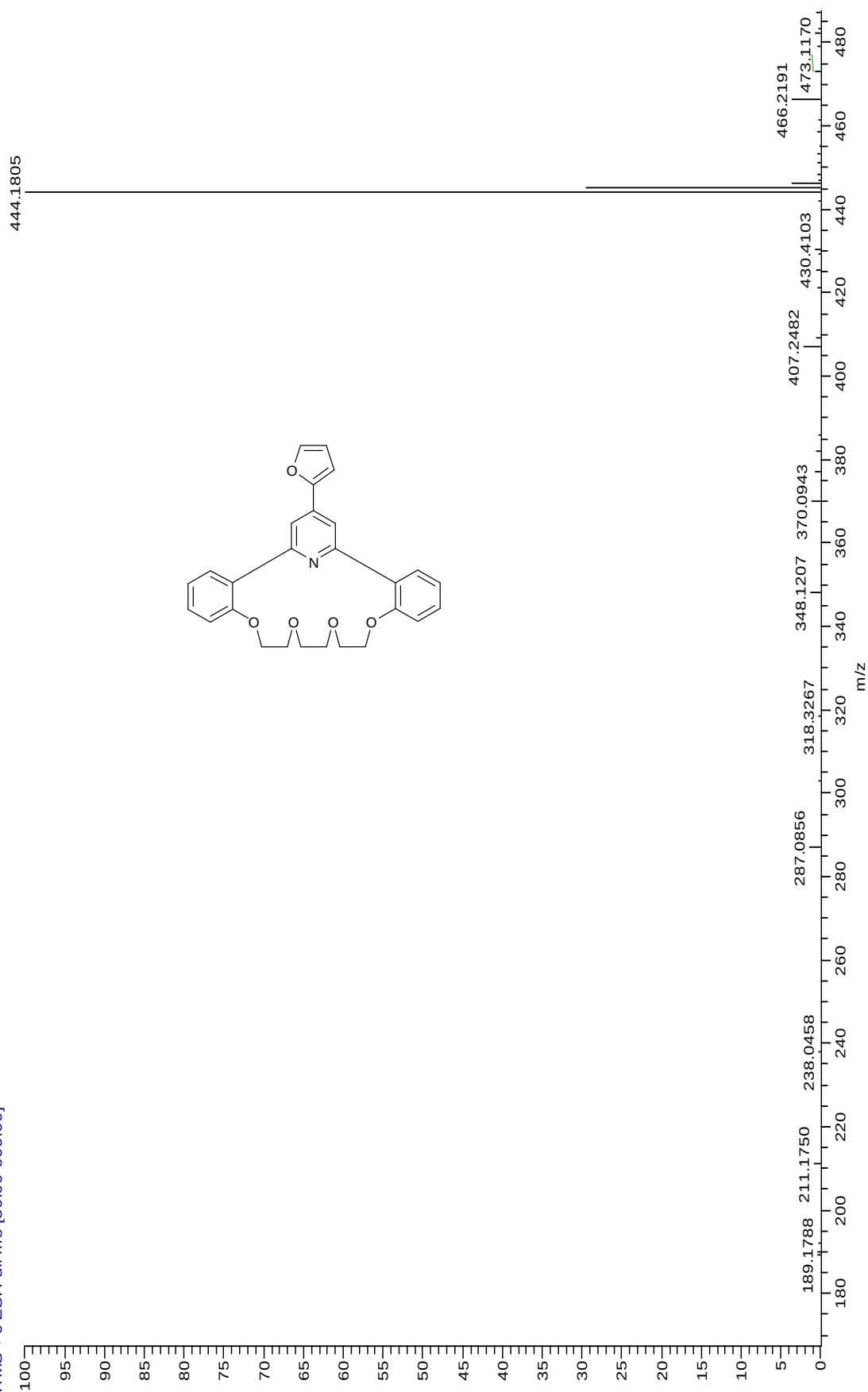
Department of Inorganic Chemistry, HUS-VNU

5/11/2016 1:14:50 AM
Instrument: FTIR Affinity - 1S

Comment:
 File name: C:\Program Files\LabSolutions\IR\Data\Up\IR - Data\DH KHTN\Tung\Tung 20 x11.ispd
 No of Scans: 20
 Intensity Mode: % Transmittance
 Min: 400 cm⁻¹
 Resolution: 4 [cm⁻¹]
 Atmosphere Correction: OFF
 Max: 4000 cm⁻¹

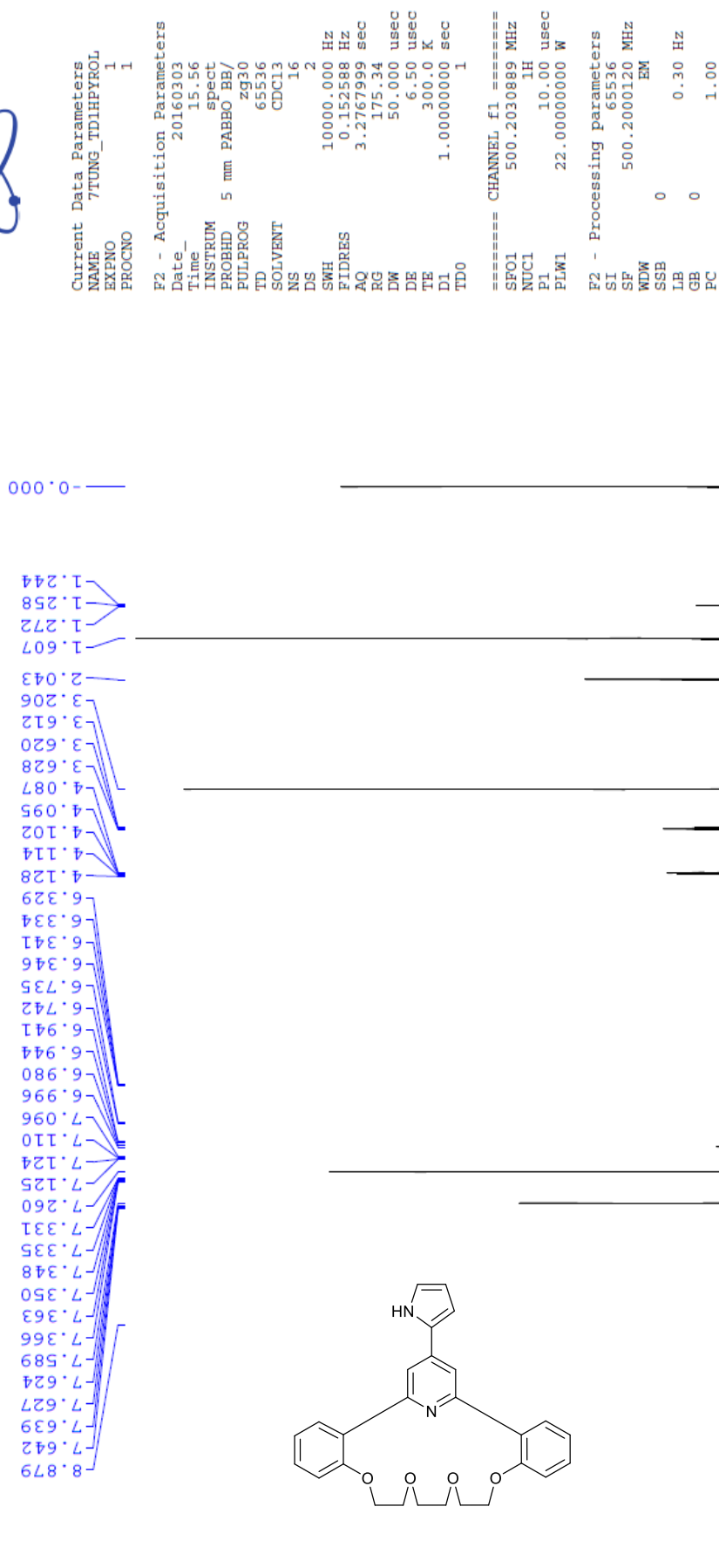
C:\LTQ Orbitrap\...TA_m2_160304104247 3/29/2016 2:23:55 P.M.
MEOH

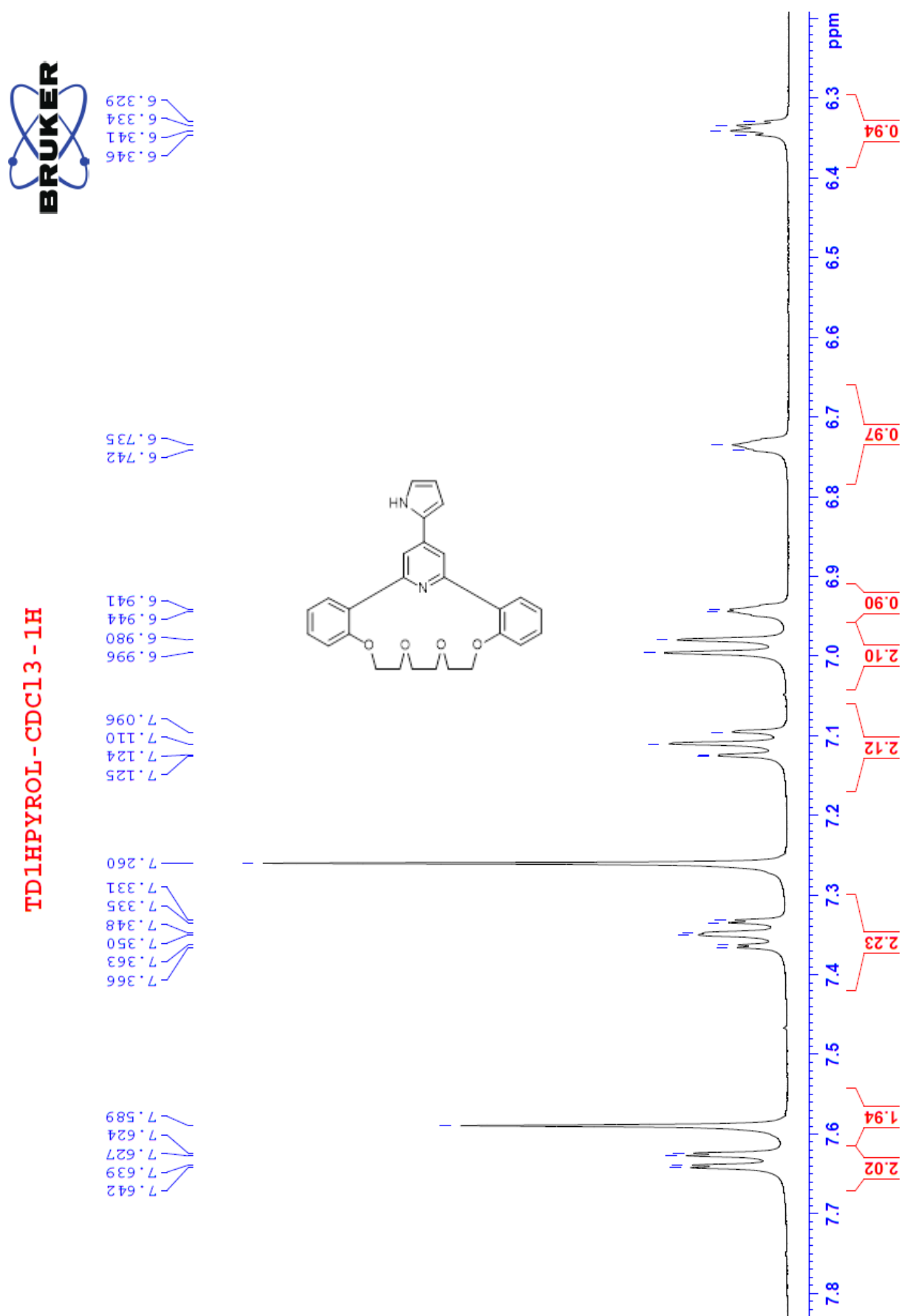
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T: ITMS+ c ESI Full ms [50.00-600.00]

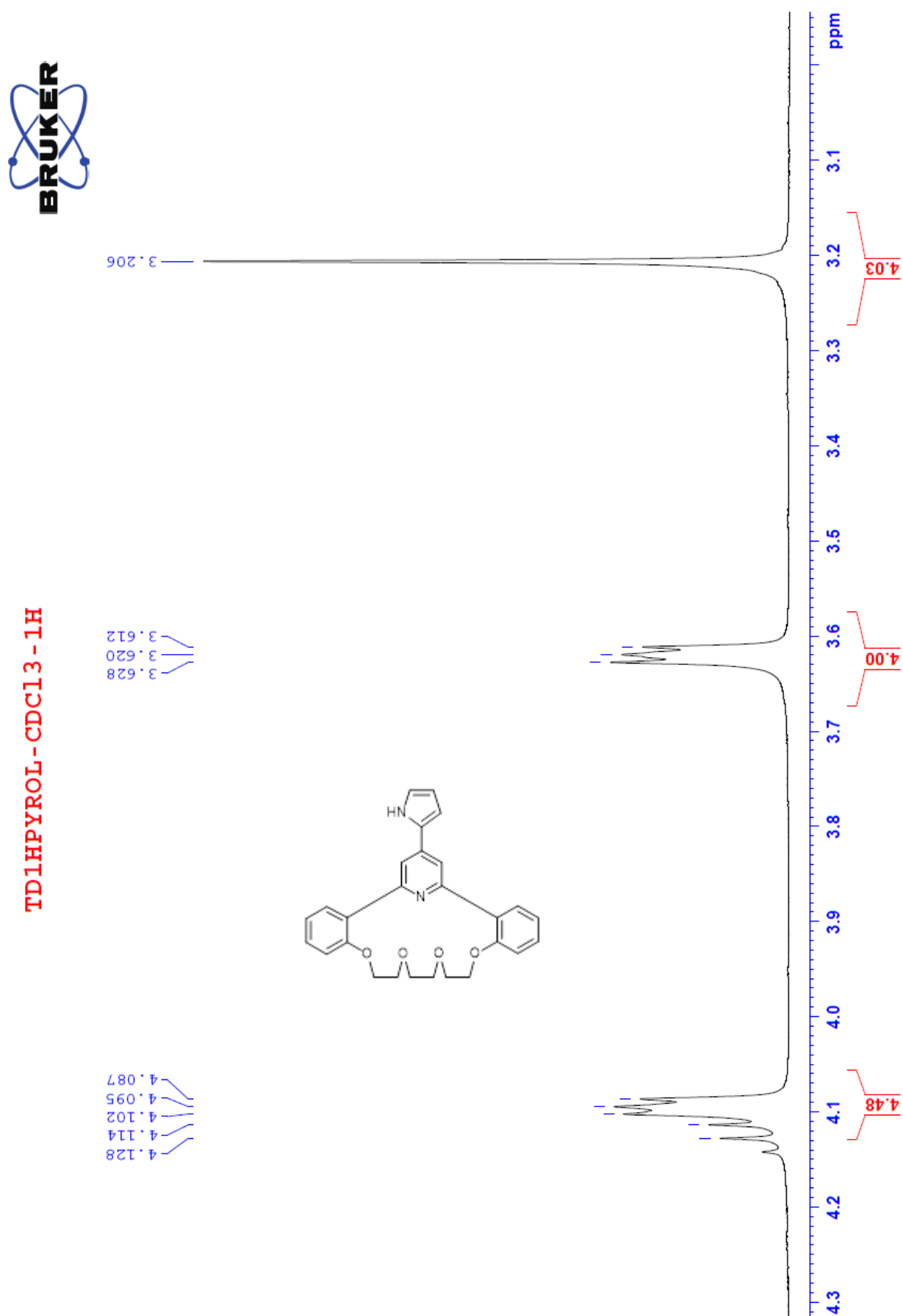




TD1HPYROL-CDCl3-1H









TD1HPYROL-CDCl₃-Cl₃CPD

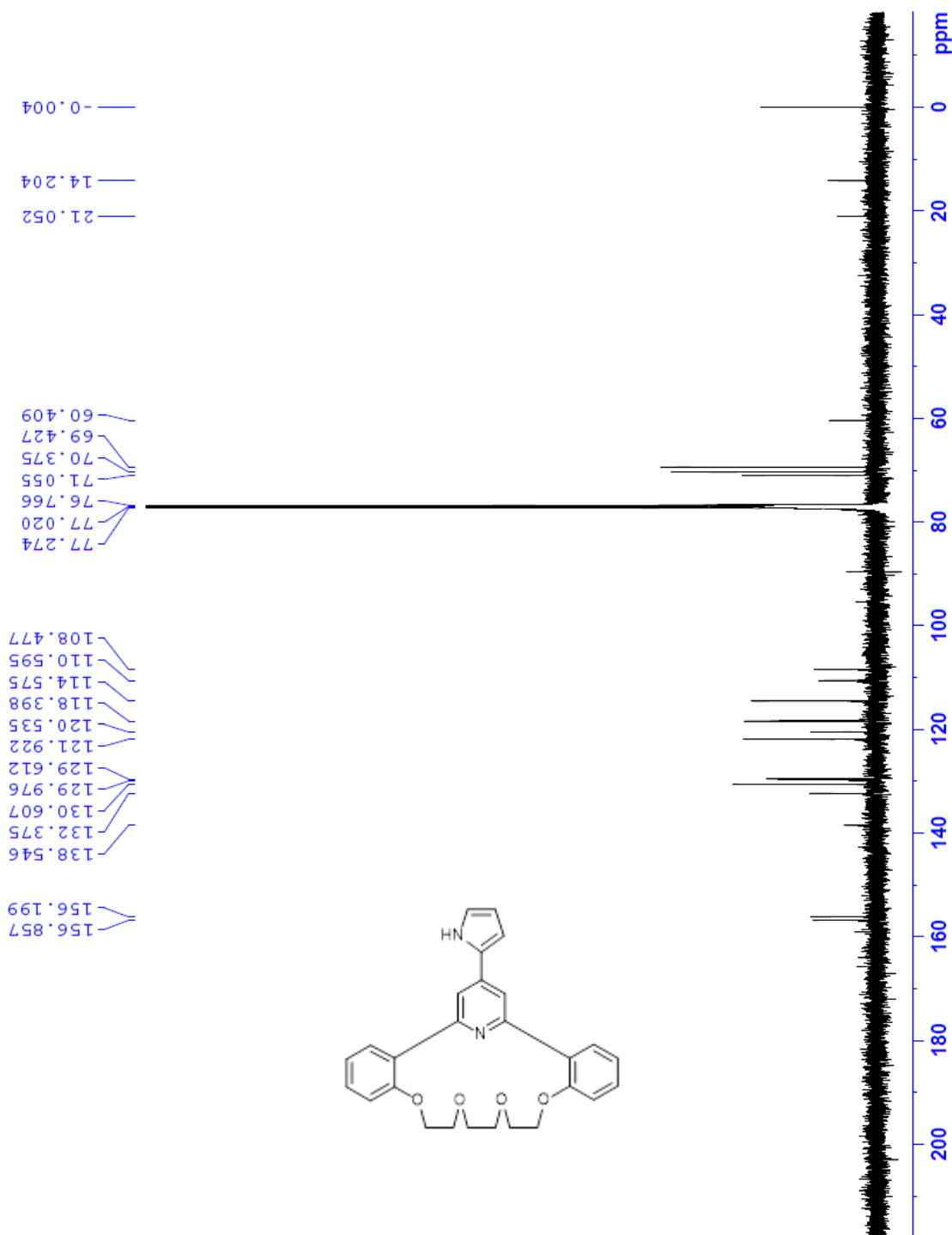
Current Data Parameters
 NAME 7TUNG_TD1HPYROL
 EXFNO 2
 PROCNO 1

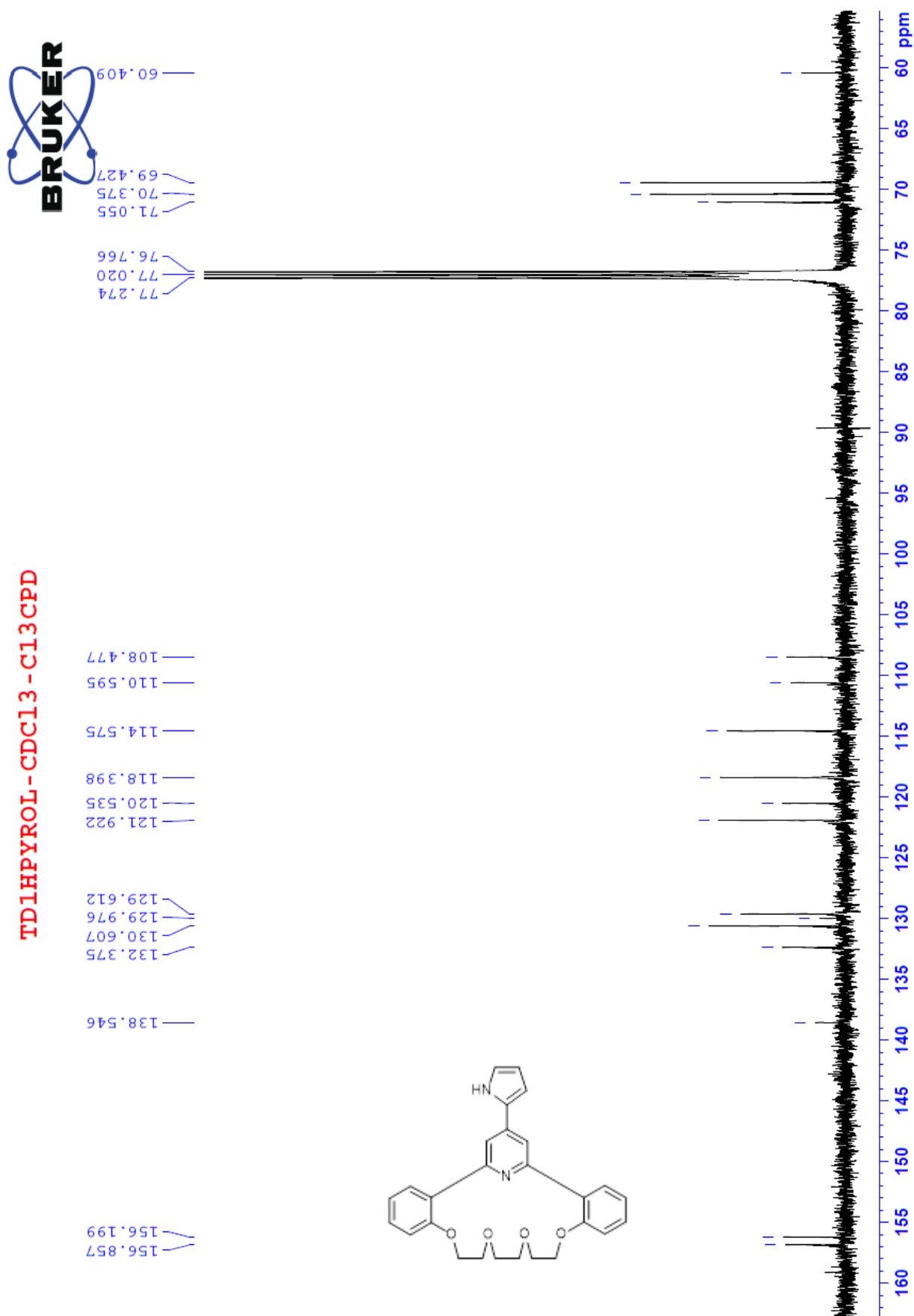
F2 - Acquisition Parameters
 Date_ 20160308
 Time 18.44
 INSTRUM spect
 PROBHD 5 mm PABBO BB/
 PULPROG zgpg30
 TD 65536
 SOLVENT CDCl₃
 NS 2048
 DS 4
 SWH 29761.904 Hz
 FIDRES 0.454131 Hz
 AQ 1.1010048 sec
 RG 198.57
 DW 16.800 usec
 DE 6.50 usec
 TE 300.5 K
 D1 2.00000000 sec
 D11 0.03000000 sec
 TD0 1

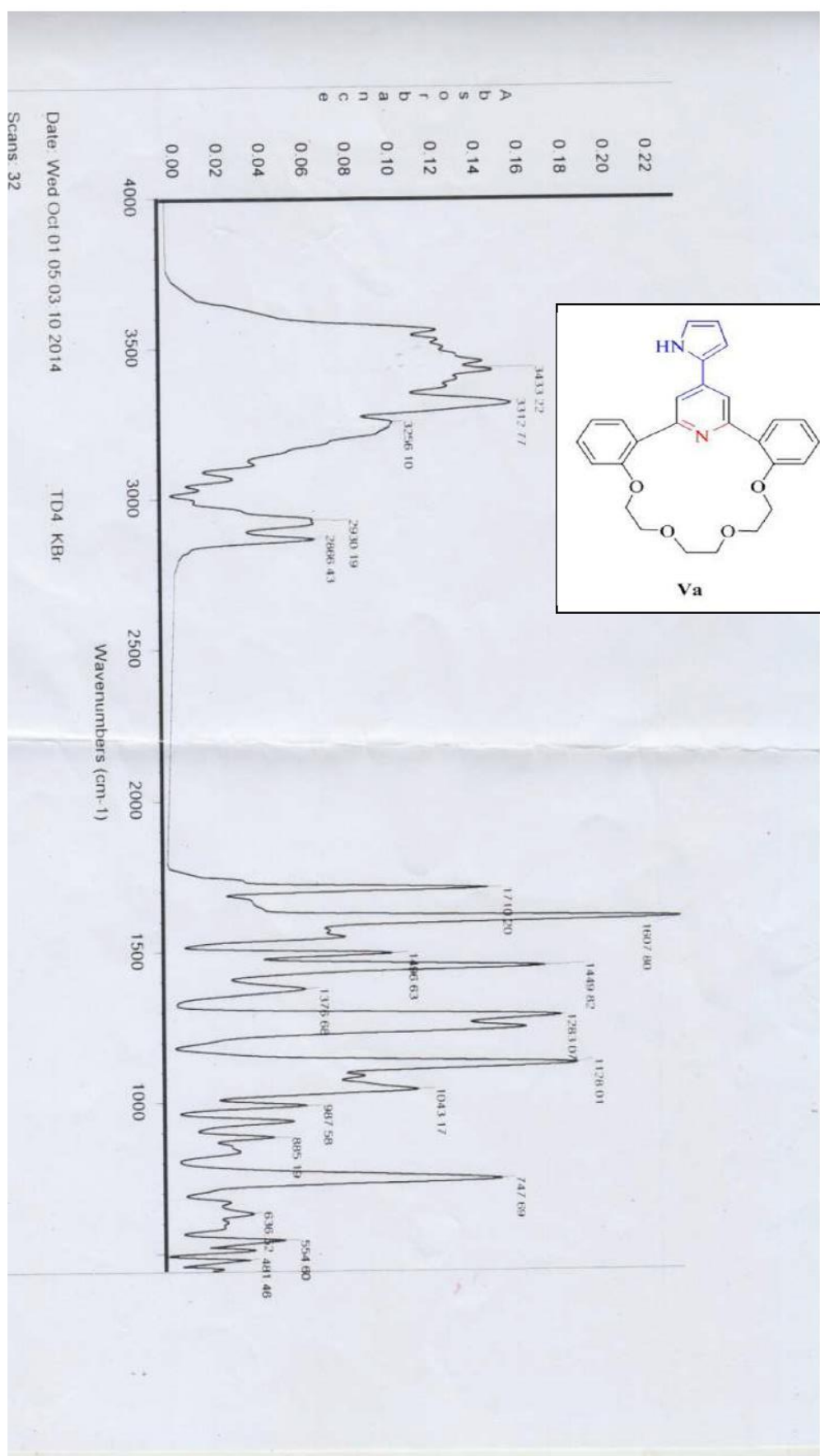
===== CHANNEL f1 =====
 SFO1 125.7879670 MHz
 NUC1 ¹³C
 P1 10.00 usec
 PLW1 88.00000000 W

===== CHANNEL f2 =====
 SFO2 500.2020008 MHz
 NUC2 ¹H
 CPDPRG2 waltz16
 PCPD2 80.00 usec
 PLW2 22.00000000 W
 PLW12 0.34375000 W
 PLW13 0.22000000 W

F2 - Processing parameters
 SI 32768
 SF 125.7753894 MHz
 EM
 WDW 0
 SSB 0
 LB 1.00 Hz
 GB 0
 PC 1.40







C:\LTQ Orbitrap\ Nhung\ Nam 2015\T3\IP4 3/4/2015 3:19:20 PM

P4 #20 RT: 0.22 AV: 1 NL: 3.52E4
T: ITMS + c ESI Full ms [150.00-2000.00]