

Supporting Information

The diversity of alkylation/acylation products of uracil and its derivatives: synthesis and structural study

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Table S1 Input data for the regression lines using NMR signals of the Table 2 and the corresponding DFT calculations.

Code of the molecule	Nucleus	Location in the molecule	NMR signal	DFT predicted signal
1	N	N1	-243.5	-261.1
1	H	H1	10.48	6.60
1	C	C2	151.6	155.5
1	N	N3	-225.0	-243.1
1	H	H3	9.58	6.52

1	C	C4	164.3	167.9
1	C	C5	99.1	103.7
1	H	H5	5.21	5.20
1	C	C6	152.4	163.4
1	H	H6	.	.
1	C	CH ₃ atC5	.	.
1	C	CH ₃ atC6	18.4	21.76
1	H	HMeatC6	1.94	2.14
1	C	COOCMe ₃ atN1	.	.
1	C	COOCMe ₃ atN3	.	.
1	C	CMe ₃ COOat N1	.	.
1	C	CMe ₃ COOat N3	.	.
1	H	HMe ₃ atN1	.	.
1	H	HMe ₃ atN1	.	.
2	N	N1	-252.2	-269.7
2	H	H1	10.16	6.58
2	C	C2	151.4	155.3
2	N	N3	-223.5	-241.3
2	H	H3	9.74	6.70
2	C	C4	164.9	169.5
2	C	C5	111.7	116.4
2	H	H5	.	.
2	C	C6	136.5	145.2
2	H	H6	6.74	7.22
2	C	CH ₃ atC5	11.6	15.9
2	H	HmeatC5	1.61	1.81
2	C	CH ₃ atC6	.	.
2	H	HmeatC6	.	.
2	C	COOCMe ₃ atN1	.	.
2	C	COOCMe ₃ atN3	.	.
2	C	CMe ₃ COOat N1	.	.
2	C	CMe ₃ COOat N3	.	.
2	H	HMe ₃ atN1	.	.
2	H	HMe ₃ atN1	.	.
3	N	N1	-249.7	-266.6
3	H	H1	10.40	6.74
3	C	C2	151.3	155.5
3	N	N3	-221.3	-239.5
3	H	H3	10.16	6.65
3	C	C4	164.3	168.2
3	C	C5	100.4	104.9
3	H	H5	5.28	5.36
3	C	C6	140.6	149.7
3	H	H6	6.87	7.39
3	C	CH ₃ atC5	.	.
3	H	HMeatC5	.	.

3	C	CH ₃ atC6	.	.
3	H	HMeatC6	.	.
3	C	COOCMe ₃ atN1	.	.
3	C	COOCMe ₃ atN3	.	.
3	C	CMe ₃ COOatN1	.	.
3	C	CMe ₃ COOatN3	.	.
3	H	HMe ₃ atN1	.	.
3	H	HMe ₃ atN1	.	.
1a	N	N1	-215.4	-221.5
1a	H	H1	.	.
1a	C	C2	148.9	157.2
1a	N	N3	-227.3	-241.6
1a	H	H3	8.73	6.87
1a	C	C4	162.2	166.4
1a	C	C5	101.6	107.2
1a	H	H5	5.55	5.45
1a	C	C6	149.7	163.5
1a	H	H6	.	.
1a	C	CH ₃ atC5	.	.
1a	H	HMeatC5	.	.
1a	C	CH ₃ atC6	18.6	22.4
1a	H	HMeatC6	2.21	2.15
1a	C	COOCMe ₃ atN1	148.2	155.7
1a	C	COOCMe ₃ atN3	.	.
1a	C	CMe ₃ COOatN1	87.7	104.1
1a	C	CMe ₃ COOatN3	.	.
1a	C	CH ₃ CCOOatN1	27.4	30.5
1a	C	CH ₃ CCOOatN3	.	.
1a	H	HMe ₃ atN1	1.60	1.53
1a	H	HMe ₃ atN3	.	.
2a	N	N1	-225.6	-228.9
2a	H	H1	.	.
2a	C	C2	148.5	153.7
2a	N	N3	-225.1	-240.7
2a	H	H3	8.67	6.75
2a	C	C4	163.3	168.6
2a	C	C5	112.0	119.4
2a	H	H5	.	.
2a	C	C6	135.3	144.6
2a	H	H6	7.71	7.62
2a	C	CH ₃ atC5	12.4	16.2
2a	H	HMeatC5	1.96	1.87
2a	C	CH ₃ atC6	.	.
2a	H	HMeatC6	.	.
2a	C	COOCMe ₃ atN1	147.3	159.4
2a	C	COOCMe ₃ atN3	.	.

2a	C	CMe ₃ COOatN1	86.9	97.5
2a	C	CMe ₃ COOatN3	.	.
2a	C	CH ₃ CCOOatN1	27.8	28.7
2a	C	CH ₃ CCOOatN3	.	.
2a	H	HMe ₃ atN1	1.61	1.63
2a	H	HMe ₃ atN3	.	.
3a	N	N1	-223.9	-226.9
3a	H	H1	.	.
3a	C	C2	147.1	153.6
3a	N	N3	-223.6	-239.2
3a	H	H3	8.76	6.72
3a	C	C4	162.5	167.2
3a	C	C5	103.7	107.9
3a	H	H5	5.80	5.57
3a	C	C6	139.7	149.3
3a	H	H6	7.88	7.80
3a	C	CH ₃ atC5	.	.
3a	H	HMeatC5	.	.
3a	C	CH ₃ atC6	.	.
3a	H	HMeatC6	.	.
3a	C	COOCMe ₃ atN1	148.1	159.1
3a	C	COOCMe ₃ atN3	.	.
3a	C	CMe ₃ COOatN1	87.4	98.0
3a	C	CMe ₃ COOatN3	.	.
3a	C	CH ₃ CCOOatN1	27.8	28.7
3a	C	CH ₃ CCOOatN3	.	.
3a	H	HMe ₃ atN1	1.61	1.65
3a	H	HMe ₃ atN3	.	.
1b	N	N1	-246.8	-261.0
1b	H	H1	9.82	6.80
1b	C	C2	150.5	155.7
1b	N	N3	-193.1	-202.1
1b	H	H3	.	.
1b	C	C4	160.8	168.4
1b	C	C5	100.3	103.9
1b	H	H5	5.56	5.37
1b	C	C6	151.3	164.0
1b	H	H6	.	.
1b	C	CH ₃ atC5	.	.
1b	H	HMeatC5	.	.
1b	C	CH ₃ atC6	18.9	21.9
1b	H	HMeatC6	2.14	2.17
1b	C	COOCMe ₃ atN1	.	.
1b	C	COOCMe ₃ atN3	147.6	159.8
1b	C	CMe ₃ COOatN1	.	.
1b	C	CMe ₃ COOatN3	86.7	97.2

1b	C	CH ₃ CCOOatN1	.	.
1b	C	CH ₃ CCOOatN3	27.4	28.3
1b	H	HMe ₃ atN1	.	.
1b	H	HMe ₃ atN3	1.60	1.59
2b	N	N1	−255.6	−269.5
2b	H	H1	9.95	6.78
2b	C	C2	150.7	155.6
2b	N	N3	−191.6	−200.0
2b	H	H3	.	.
2b	C	C4	161.9	170.0
2b	C	C5	110.5	116.9
2b	H	H5	.	.
2b	C	C6	135.9	145.6
2b	H	H6	7.04	7.26
2b	C	CH ₃ atC5	12.3	15.9
2b	H	HMeatC5	1.91	1.86
2b	C	CH ₃ atC6	.	.
2b	H	HMeatC6	.	.
2b	C	COOCMe ₃ atN1	.	.
2b	C	COOCMe ₃ atN3	147.8	159.7
2b	C	CMe ₃ COOatN1	.	.
2b	C	CMe ₃ COOatN3	87.0	97.4
2b	C	CH ₃ CCOOatN1	.	.
2b	C	CH ₃ CCOOatN3	27.5	28.3
2b	H	HMe ₃ atN1	.	.
2b	H	HMe ₃ atN3	1.60	1.60
3b	N	N1	−253.4	−266.4
3b	H	H1	10.04	6.92
3b	C	C2	150.5	155.9
3b	N	N3	−189.7	−198.1
3b	H	H3	.	.
3b	C	C4	161.0	168.9
3b	C	C5	102.0	105.3
3b	H	H5	5.76	5.52
3b	C	C6	140.1	149.9
3b	H	H6	.	7.43
3b	C	CH ₃ atC5	.	.
3b	H	HMeatC5	.	.
3b	C	CH ₃ atC6	.	.
3b	H	HMeatC6	.	.
3b	C	COOCMe ₃ atN1	.	.
3b	C	COOCMe ₃ atN3	147.6	159.6
3b	C	CMe ₃ COOatN1	.	.
3b	C	CMe ₃ COOatN3	87.1	97.5
3b	C	CH ₃ CCOOatN1	.	.
3b	C	CH ₃ CCOOatN3	27.5	28.3

3b	H	HMe ₃ atN1	.	.
3b	H	HMe ₃ atN3	1.61	1.60
1c	N	N1	-216.4	-221.0
1c	H	H1	.	.
1c	C	C2	147.8	156.4
1c	N	N3	-194.0	-201.6
1c	H	H3	.	.
1c	C	C4	159.5	166.9
1c	C	C5	101.2	106.2
1c	H	H5	5.54	5.52
1c	C	C6	148.9	163.1
1c	H	H6	.	.
1c	C	CH ₃ atC5	.	.
1c	H	HMeatC5	.	.
1c	C	CH ₃ atC6	18.6	21.7
1c	H	HMeatC6	2.18	2.20
1c	C	COOCMe ₃ atN1	147.3	158.6
1c	C	COOCMe ₃ atN3	147.1	158.8
1c	C	CMe ₃ COOatN1	88.1	99.8
1c	C	CMe ₃ COOatN3	87.1	98.0
1c	C	CH ₃ CCOOatN1	27.4	28.3
1c	C	CH ₃ CCOOatN3	27.3	28.3
1c	H	HMe ₃ atN1	1.60	1.63
1c	H	HMe ₃ atN3	1.59	1.60
2c	N	N1	-226.5	-229.0
2c	H	H1	.	.
2c	C	C2	145.7	154.2
2c	N	N3	.	-199.3
2c	H	H3	.	.
2c	C	C4	161.1	169.3
2c	C	C5	111.7	119.6
2c	H	H5	.	.
2c	C	C6	134.6	144.8
2c	H	H6	7.68	7.68
2c	C	CH ₃ atC5	12.5	16.2
2c	H	HMeatC5	1.95	1.92
2c	C	CH ₃ atC6	.	.
2c	H	HMeatC6	.	.
2c	C	COOCMe ₃ atN1	148.4	158.9
2c	C	COOCMe ₃ atN3	147.4	158.8
2c	C	CMe ₃ COOatN1	87.3	98.5
2c	C	CMe ₃ COOatN3	87.0	98.1
2c	C	CH ₃ CCOOatN1	27.8	28.7
2c	C	CH ₃ CCOOatN3	27.4	28.3
2c	H	HMe ₃ atN1	1.59	1.65
2c	H	HMe ₃ atN3	1.59	1.61

3c	N	N1	-224.8	-227.2
3c	H	H1	.	.
3c	C	C2	148.0	154.2
3c	N	N3	-190.0	-197.8
3c	H	H3	.	.
3c	C	C4	159.9	168.0
3c	C	C5	103.4	108.0
3c	H	H5	5.81	5.72
3c	C	C6	138.9	149.3
3c	H	H6	7.87	7.90
3c	C	CH ₃ atC5	.	.
3c	H	HMeatC5	.	.
3c	C	CH ₃ atC6	.	.
3c	H	HMeatC6	.	.
3c	C	COOCMe ₃ atN1	147.2	158.3
3c	C	COOCMe ₃ atN3	145.5	158.8
3c	C	CMe ₃ COOatN1	87.1	99.0
3c	C	CMe ₃ COOatN3	87.6	98.2
3c	C	CH ₃ CCOOatN1	27.7	28.7
3c	C	CH ₃ CCOOatN3	27.4	28.3
3c	H	HMe ₃ atN1	1.61	1.66
3c	H	HMe ₃ atN3	1.60	1.60
1d	N	N1	-114.8	-146.8
1d	H	H1	.	.
1d	C	C2	160.0	170.6
1d	N	N3	-136.3	-169.0
1d	H	H3	.	.
1d	C	C4	166.7	177.4
1d	C	C5	108.7	110.6
1d	H	H5	7.01	6.68
1d	C	C6	173.4	184.1
1d	H	H6	.	.
1d	C	CH ₃ atC5	.	.
1d	H	HMeatC5	.	.
1d	C	CH ₃ atC6	24.1	26.5
1d	H	HMeatC6	2.56	2.52
1d	C	COOCMe ₃ atO2	149.4	157.5
1d	C	COOCMe ₃ atO4	148.7	156.3
1d	C	CMe ₃ COOatO2	85.4	97.5
1d	C	CMe ₃ COOatO4	84.7	97.9
1d	C	CH ₃ CCOOatO2	27.6	28.3
1d	C	CH ₃ CCOOatO4	27.5	28.2
1d	H	HMe ₃ atO2	1.54	1.64
1d	H	HMe ₃ atO4	1.55	1.55

Table S2 Input data for the regression lines using NMR signals of the Table 4 and the corresponding DFT calculations.

Code of the molecule	Nucleus	Location in the molecule	NMR signal	DFT predicted signal
1	N	N1	−243.5	−261.1
1	H	H1	10.48	6.60
1	C	C2	151.6	155.5
1	N	N3	−225.0	−243.1
1	H	H3	9.58	6.52
1	C	C4	164.3	167.9
1	C	C5	99.1	103.7
1	H	H5	5.21	5.20
1	C	C6	152.4	163.4
1	H	H6	.	.
1	C	CH ₃ atC5	.	.
1	C	CH ₃ atC6	18.4	21.76
1	H	HMeatC6	1.94	2.14
1	C	COOCMe ₃ atN1	.	.
1	C	COOCMe ₃ atN3	.	.
1	C	CMe ₃ COOat N1	.	.
1	C	CMe ₃ COOat N3	.	.
1	H	HMe ₃ atN1	.	.
1	H	HMe ₃ atN1	.	.
2	N	N1	−252.2	−269.7
2	H	H1	10.16	6.58
2	C	C2	151.4	155.3
2	N	N3	−223.5	−241.3
2	H	H3	9.74	6.70
2	C	C4	164.9	169.5
2	C	C5	111.7	116.4
2	H	H5	.	.
2	C	C6	136.5	145.2
2	H	H6	6.74	7.22
2	C	CH ₃ atC5	11.6	15.9
2	H	HMeatC5	1.61	1.81
2	C	CH ₃ atC6	.	.
2	H	HMeatC6	.	.
2	C	COOCMe ₃ atN1	.	.
2	C	COOCMe ₃ atN3	.	.
2	C	CMe ₃ COOat N1	.	.
2	C	CMe ₃ COOat N3	.	.
2	H	HMe ₃ atN1	.	.
2	H	HMe ₃ atN1	.	.
3	N	N1	−249.7	−266.6
3	H	H1	10.40	6.74
3	C	C2	151.3	155.5

3	N	N3	-221.3	-239.5
3	H	H3	10.16	6.65
3	C	C4	164.3	168.2
3	C	C5	100.4	104.9
3	H	H5	5.28	5.36
3	C	C6	140.6	149.7
3	H	H6	6.87	7.39
3	C	CH ₃ atC5	.	.
3	H	HMeatC5	.	.
3	C	CH ₃ atC6	.	.
3	H	HMeatC6	.	.
3	C	COOCMe ₃ atN1	.	.
3	C	COOCMe ₃ atN3	.	.
3	C	CMe ₃ COOatN1	.	.
3	C	CMe ₃ COOatN3	.	.
3	H	HMe ₃ atN1	.	.
3	H	HMe ₃ atN1	.	.
1i	N	N1	-246.0	-261.5
1i	H	H1	9.83	6.72
1i	C	C2	152.9	157.4
1i	N	N3	-213.4	-225.1
1i	H	H3	.	.
1i	C	C4	163.1	169.3
1i	C	C5	100.4	103.8
1i	H	H5	5.56	5.31
1i	C	C6	149.4	160.9
1i	H	H6	.	.
1i	C	CH ₃ atC5	.	.
1i	H	HMeatC5	.	.
1i	C	CH ₃ atC6	18.7	21.4
1i	H	HMeatC6	2.15	2.13
1i	C	CH ₂ MeatN1	.	.
1i	C	CH ₂ MeatN3	35.5	39.9
1i	H	HCH ₂ MeatN1	.	.
1i	H	HCH ₂ MeatN3	3.96	3.71
1i	C	CH ₃ CH ₂ atN1	.	.
1i	C	CH ₃ CH ₂ atN3	12.9	14.8
1i	H	HMeatN1	.	.
1i	H	HMeatN3	1.22	1.12
1j	N	N1	-237.7	-244.1
1j	H	H1	.	.
1j	C	C2	151.6	159.8
1j	N	N3	-212.8	-223.7
1j	H	H3	.	.
1j	C	C4	162.1	167.8
1j	C	C5	101.7	105.4

1j	H	H5	5.55	5.40
1j	C	C6	150.9	163.8
1j	H	H6	.	.
1j	C	CH ₃ atC5	.	.
1j	H	HMeatC5	.	.
1j	C	CH ₃ atC6	19.5	22.3
1j	H	HMeatC6	2.22	2.23
1j	C	CH ₂ MeatN1	40.1	45.4
1j	C	CH ₂ MeatN3	36.3	40.8
1j	H	HCH ₂ MeatN1	3.87	3.82
1j	H	HCH ₂ MeatN3	3.96	3.76
1j	C	CH ₃ CH ₂ atN1	14.1	15.5
1j	C	CH ₃ CH ₂ atN3	12.8	14.6
1j	H	HMeatN1	1.26	1.22
1j	H	HMeatN3	1.19	1.14
1l	N	N1	-179.0	-194.9
1l	H	H1	.	.
1l	C	C2	155.6	164.9
1l	N	N3	-215.8	-226.2
1l	H	H3	.	.
1l	C	C4	163.1	167.9
1l	C	C5	105.9	110.3
1l	H	H5	5.95	5.80
1l	C	C6	162.5	172.7
1l	H	H6	.	.
1l	C	CH ₃ atC5	.	.
1l	H	HMeatC5	.	.
1l	C	CH ₃ atC6	23.6	26.4
1l	H	HMeatC6	2.16	2.19
1l	C	CH ₂ MeatO2	64.7	71.6
1l	C	CH ₂ MeatN3	36.1	40.9
1l	H	HCH ₂ MeatO2	4.45	4.34
1l	H	HCH ₂ MeatN3	4.03	3.84
1l	C	CH ₃ CH ₂ atO2	14.2	16.3
1l	C	CH ₃ CH ₂ atN3	13.3	15.1
1l	H	HMeatO2	1.41	1.51
1l	H	HMeatN3	1.23	1.19
2h	N	N1	-244.3	-249.9
2h	H	H1	.	.
2h	C	C2	150.8	156.9
2h	N	N3	-225.0	-241.1
2h	H	H3	9.17	6.81
2h	C	C4	164.3	169.1
2h	C	C5	110.9	116.8
2h	H	H5	.	.
2h	C	C6	140.0	149.6

2h	H	H6	7.00	7.29
2h	C	CH ₃ atC5	12.4	16.1
2h	H	HMeatC5	1.93	1.83
2h	C	CH ₃ atC6	.	.
2h	H	HMeatC6	.	.
2h	C	CH ₂ MeatN1	43.6	49.2
2h	C	CH ₂ MeatN3	.	.
2h	H	HCH ₂ MeatN1	3.78	3.67
2h	H	HCH ₂ MeatN3	.	.
2h	C	CH ₃ CH ₂ atN1	14.4	16.8
2h	C	CH ₃ CH ₂ atN3	.	.
2h	H	HMeatN1	1.29	1.21
2h	H	HMeatN3	.	.
2j	N	N1	-245.5	-250.9
2j	H	H1	.	.
2j	C	C2	151.0	158.7
2j	N	N3	-212.1	-222.7
2j	H	H3	.	.
2j	C	C4	163.6	170.3
2j	C	C5	109.9	116.2
2j	H	H5	.	.
2j	C	C6	137.8	147.7
2j	H	H6	6.98	7.30
2j	C	CH ₃ atC5	13.0	16.8
2j	H	HMeatC5	1.93	1.86
2j	C	CH ₃ atC6	.	.
2j	H	HMeatC6	.	.
2j	C	CH ₂ MeatN1	44.4	50.1
2j	C	CH ₂ MeatN3	36.5	40.9
2j	H	HCH ₂ MeatN1	3.78	3.72
2j	H	HCH ₂ MeatN3	4.03	3.83
2j	C	CH ₃ CH ₂ atN1	14.3	16.7
2j	C	CH ₃ CH ₂ atN3	12.8	14.7
2j	H	HMeatN1	1.31	1.23
2j	H	HMeatN3	1.22	1.15
3h	N	N1	-241.2	-246.5
3h	H	H1	.	.
3h	C	C2	150.9	157.2
3h	N	N3	-222.9	-239.2
3h	H	H3	.	6.77
3h	C	C4	164.1	167.8
3h	C	C5	102.4	105.2
3h	H	H5	5.72	5.38
3h	C	C6	144.0	153.9
3h	H	H6	7.19	7.47
3h	C	CH ₃ atC5	.	.

3h	H	HMeatC5	.	.
3h	C	CH ₃ atC6	.	.
3h	H	HMeatC6	.	.
3h	C	CH ₂ MeatN1	43.9	49.9
3h	C	CH ₂ MeatN3	.	.
3h	H	HCH ₂ MeatN1	3.80	3.70
3h	H	HCH ₂ MeatN3	.	.
3h	C	CH ₃ CH ₂ atN1	14.4	16.8
3h	C	CH ₃ CH ₂ atN3	.	.
3h	H	HMeatN1	1.32	1.22
3h	H	HMeatN3	.	.
3i	N	N1	-253.4	-267.0
3i	H	H1	9.78	6.83
3i	C	C2	152.8	157.5
3i	N	N3	-209.8	-220.6
3i	H	H3	.	.
3i	C	C4	163.0	169.7
3i	C	C5	102.3	105.1
3i	H	H5	5.77	5.45
3i	C	C6	138.1	147.4
3i	H	H6	7.16	7.35
3i	C	CH ₃ atC5	.	.
3i	H	HMeatC5	.	.
3i	C	CH ₃ atC6	.	.
3i	H	HMeatC6	.	.
3i	C	CH ₂ MeatN1	.	.
3i	C	CH ₂ MeatN3	35.8	40.2
3i	H	HCH ₂ MeatN1	.	.
3i	H	HCH ₂ MeatN3	3.99	3.74
3i	C	CH ₃ CH ₂ atN1	.	.
3i	C	CH ₃ CH ₂ atN3	12.8	14.8
3i	H	HMeatN1	.	.
3i	H	HMeatN3	1.24	1.14
3j	N	N1	-242.3	-247.3
3j	H	H1	.	.
3j	C	C2	151.0	159.2
3j	N	N3	-210.0	-220.6
3j	H	H3	.	.
3j	C	C4	163.0	169.2
3j	C	C5	101.8	105.0
3j	H	H5	5.70	5.48
3j	C	C6	141.7	151.6
3j	H	H6	7.13	7.43
3j	C	CH ₃ atC5	.	.
3j	H	HMeatC5	.	.
3j	C	CH ₃ atC6	.	.

3j	H	HMeatC6	.	.
3j	C	CH ₂ MeatN1	44.7	50.9
3j	C	CH ₂ MeatN3	36.2	40.8
3j	H	HCH ₂ MeatN1	3.79	3.73
3j	H	HCH ₂ MeatN3	3.98	3.77
3j	C	CH ₃ CH ₂ atN1	14.3	16.6
3j	C	CH ₃ CH ₂ atN3	12.8	14.7
3j	H	HMeatN1	1.30	1.25
3j	H	HMeatN3	1.20	1.15
3m	N	N1	-224.4	-224.3
3m	H	H1	.	.
3m	C	C2	156.5	161.6
3m	N	N3	.	-169.2
3m	H	H3	.	.
3m	C	C4	171.3	179.7
3m	C	C5	95.8	92.5
3m	H	H5	5.85	5.78
3m	C	C6	146.1	158.3
3m	H	H6	7.40	7.83
3m	C	CH ₃ atC5	.	.
3m	H	HMeatC5	.	.
3m	C	CH ₃ atC6	.	.
3m	H	HMeatC6	.	.
3m	C	CH ₂ MeatN1	45.4	51.5
3m	C	CH ₂ MeatO4	63.0	69.9
3m	H	HCH ₂ MeatN1	3.90	3.76
3m	H	HCH ₂ MeatO4	4.43	4.42
3m	C	CH ₃ CH ₂ atN1	14.5	16.7
3m	C	CH ₃ CH ₂ atO4	14.2	16.2
3m	H	HMeatN1	1.35	1.24
3m	H	HMeatO4	1.36	1.31

Table S3 Linear regression parameters ¹for the correspondence between the NMR chemical shifts² and the DFT³ predicted chemical shifts⁴ in ppm.

NMR source data	Regression line	<i>n</i>	<i>R</i> ²
Table 2 (Boc ₂ O)			
¹⁵ N	<i>NMR</i> = 23.23 + 1.046 <i>DFT</i>	25	0.9454
¹ H	<i>NMR</i> = -0.4812 + 1.2667 <i>DFT</i>	51	0.8807
¹³ C	<i>NMR</i> = -1.986 + 0.9559 <i>DFT</i>	103	0.9962
Table 4 (EtI)			
¹⁵ N	<i>NMR</i> = -1.791 + 0.9426 <i>DFT</i>	23	0.9181
¹ H	<i>NMR</i> = -2.427 + 0.9716 <i>DFT</i>	83	0.9986
¹³ C	<i>NMR</i> = -0.3466 + 1.2027 <i>DFT</i>	61	0.8657

¹*n* is the number of data included, *R*² is the determination coefficient

²NMR data are given in the Table 2 and Table 4

³ DFT data are given in the Table S8A and Table S8B

⁴ Based on the calculated linear regression line the lacking value for the ¹⁵N chemical shift for the *N*3 nitrogen atom in the **2c** compound was – 185 ppm with the 95 % confidence interval of [–202;–168] ppm and for the *N*3 nitrogen atom in the **3m** compound was –161 ppm with the 95 % confidence interval of [–176; –147] ppm.

Table S4 (Boc)₂O reaction with uracil (**3**), thymine (**2**) and 6-methyluracil (**3**)¹. The B3LYP/6–311G(d) theoretical estimation of Δ*G* including acetonitrile as solvent.

Reaction product	Δ <i>G</i> Uracil (3) kcal/mole (kJ/mol)	Δ <i>G</i> Thymine (2) kcal/mole (kJ/mol)	Δ <i>G</i> 6-methyluracil (1) kcal/mole (kJ/mol)
<i>N</i> ¹ –Boc	–11.03 (–46.16) 3a	–10.89 (–45.58) 2a	+2.11 (+8.81) 1a
<i>O</i> ² –Boc	+2.26 (+9.45)	+3.10 (+12.96)	+2.77 (+11.59)
<i>N</i> ³ –Boc	–10.96 (–45.84) 3b	–10.13 (–42.37) 2b	–10.49 (–43.87) 1b
<i>O</i> ⁴ –Boc	–0.68 (–2.86)	+0.18 (+0.77)	–0.72 (–3.00)
<i>C</i> ⁵ –Boc	–19.18 (–80.24)	–	–15.23 (–63.71)
<i>C</i> ⁶ –Boc	–18.92 (–79.15)	–14.67 (–61.38)	–
<i>N</i> ¹ , <i>O</i> ² –diBoc	–3.98 (–16.66)	–5.49 (–22.98)	–1.45 (–6.06)
<i>N</i> ¹ , <i>N</i> ³ –diBoc	–21.08 (–88.19) 3c	–21.00 (–87.89) 2c	–18.86 (–78.91) 1c
<i>N</i> ¹ , <i>O</i> ⁴ –diBoc	–10.91 (–45.64) 3f	–10.79 (–45.17) 2f	–10.25 (–42.90) 1f
<i>N</i> ¹ , <i>C</i> ⁵ –diBoc	–29.05 (–121.57)	–	–22.73 (–95.11)
<i>N</i> ¹ , <i>C</i> ⁶ –diBoc	–24.21 (–101.30)	–22.30 (–93.29)	–
<i>O</i> ² , <i>N</i> ³ –diBoc	–3.42 (–14.32) 3e	–7.98 (–33.37) 2e	–3.40 (–14.23) 1e
<i>O</i> ² , <i>O</i> ⁴ –diBoc	–6.36 (–26.61) 3d	–6.94 (–29.03) 2d	–6.71 (–28.06) 1d
<i>O</i> ² , <i>C</i> ⁵ –diBoc	–15.83 (–66.22)	–	–13.29 (–55.61)
<i>O</i> ² , <i>C</i> ⁶ –diBoc	–16.09 (–67.33)	–12.36 (–51.70)	–
<i>N</i> ³ , <i>O</i> ⁴ –diBoc	–2.18 (–9.14)	–0.77 (–3.24)	–2.97 (–12.42)
<i>N</i> ³ , <i>C</i> ⁵ –diBoc	–29.07 (–121.62)	–	–25.36 (–106.11)
<i>N</i> ³ , <i>C</i> ⁶ –diBoc	–28.95 (–121.14)	–25.24 (–105.60)	–
<i>O</i> ⁴ , <i>C</i> ⁵ –diBoc	–11.76 (–49.19)	–	–9.42 (–39.40)
<i>O</i> ⁴ , <i>C</i> ⁶ –diBoc	–18.52 (–77.47)	–13.96 (–58.41)	–
<i>C</i> ⁵ , <i>C</i> ⁶ –diBoc	–29.85 (–124.90)	–	–

¹) Models of the (Boc)₂O reaction with **3**, **2**, **1**:



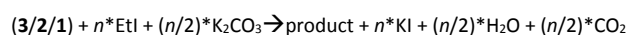
where *n*=1 for mono-substitutions, and *n*=2 for di-substitutions.

Table S5 Ethylation of uracil, thymine and 6-methyluracil ¹). The B3LYP/6–311G(d) theoretical estimation of Δ*G* including DMF as solvent.

Reaction product	Δ <i>G</i> Uracil (3) kcal/mole (kJ/mol)	Δ <i>G</i> Thymine (2) kcal/mole (kJ/mol)	Δ <i>G</i> 6-methyluracil (3) kcal/mole (kJ/mol)
<i>N</i> ¹ –Et	–39.03 (–163.32) 3h	–27.99 (–117.11) 2h	–23.73 (–99.30) 1h
<i>O</i> ² –Et	–20.85 (–87.25)	–6.77 (–28.34)	–9.20 (–38.51)
<i>N</i> ³ –Et	–38.64 (–161.69) 3i	–27.56 (–115.33) 2i	–27.20 (–113.80) 1i
<i>O</i> ⁴ –Et	–22.59 (–94.52)	–10.88 (–45.51)	–11.35 (–47.48)
<i>C</i> ⁵ –Et	–44.39 (–185.75)	–	–30.47 (–127.50)
<i>C</i> ⁶ –Et	–46.99 (–196.59)	–33.59 (–140.55)	–
<i>N</i> ¹ , <i>O</i> ² –diEt	–55.21 (–231.00)	–34.87 (–145.91)	–29.51 (–123.46)
<i>N</i> ¹ , <i>N</i> ³ –diEt	–77.39 (–323.81) 3j	–55.74 (–233.23) 2j	–52.06 (–217.83) 1j
<i>N</i> ¹ , <i>O</i> ⁴ –diEt	–57.63 (–241.13) 3m	–40.56 (–169.73) 2m	–36.24 (–151.62) 1m
<i>N</i> ¹ , <i>C</i> ⁵ –diEt	–82.88 (–346.76)	–	–53.71 (–224.71)

N^1, C^6 -diEt	-80.60 (-337.25)	-56.10 (-234.72)	-
O^2, N^3 -diEt	-59.82 (-250.27) 3l	-37.48 (-156.82) 2l	-37.93 (-158.72) 1l
O^2, O^4 -diEt	-46.35 (-193.93) 3k	-28.06 (-117.39) 2k	-24.27 (-101.53) 1k
O^2, C^5 -diEt	-61.86 (-258.81)	-	-37.90 (-158.56)
O^2, C^6 -diEt	-62.93 (-263.31)	-40.16 (-168.03)	-
N^3, O^4 -diEt	-53.68 (-224.61)	-25.50 (-106.71)	-32.24 (-134.91)
N^3, C^5 -diEt	-82.19 (-343.87)	-	-58.60 (-245.17)
N^3, C^6 -diEt	-85.03 (-355.76)	-61.25 (-256.28)	-
O^4, C^5 -diEt	-65.35 (-273.42)	-	-41.14 (-172.14)
O^4, C^6 -diEt	-69.38 (-290.28)	-44.88 (-187.78)	-
C^5, C^6 -diEt	-88.73 (-371.23)	-	-

1) Models of the EtI reaction with **1**, **2**, **3**:



where $n=1$ for mono-substitutions, and $n=2$ for di-substitutions

Table S6 Comparison of experimental yields of di-Boc and di-Et products with theoretical ones based on ΔG calculations.

	Experimental yield of the reactions ¹			Theoretical, ΔG [kcal/mol]		
	U	T	6-MeU	U	T	6-MeU
	N^1, N^3 (3c)	N^1, N^3 (2c)	N^1, N^3 (1c)	N^1, N^3	N^1, N^3	N^1, N^3
di-Boc	—	—	5% 1.5 eq D	-21.1	-21.0	-18.9
	42 % 3.0 eq D	39 % 3.0 eq D	11 % 3.0 eq D			
	77 % 3.0 eq P	89 % 3.0 eq P	20 % 3.0 eq P			
	O^2, O^4	O^2, O^4	O^2, O^4 (1d)	O^2, O^4	O^2, O^4	O^2, O^4
	—	—	12 % 1.5 eq D	-6.4	-6.9	-6.7
	—	—	32 % 3.0 eq D			
	O^2, N^3	O^2, N^3	O^2, N^3	O^2, N^3	O^2, N^3	O^2, N^3
	—	—	—	-3.4	-8.0	-3.4
	N^1, N^3 (3j)	N^1, N^3 (2j)	N^1, N^3 (1j)	N^1, N^3	N^1, N^3	N^1, N^3
	—	26 % 1.0 eq	30 % 1.0 eq	-39.0	-28.0	-23.7
di-Et	31 % 1.5 eq	30 % 1.5 eq	75 % 1.5 eq			
	29 % 3.0 eq	32 % 1.5 eq	99 % 3.0 eq			
	O^2, O^4	O^2, O^4	O^2, O^4	O^2, O^4	O^2, O^4	O^2, O^4
	—	—	—	-46.4	-28.1	-24.3
	O^2, N^3	O^2, N^3	O^2, N^3 (1l)	O^2, N^3	O^2, N^3	O^2, N^3
	—	—	3 % 1.5 eq	-59.8	-37.5	-37.9
	N^1, O^4 (3m)	N^1, O^4	N^1, O^4	N^1, O^4	N^1, O^4	N^1, O^4
	1 % 1.5 eq	—	—	-57.6	-40.6	-36.2
	1 % 3.0 eq	—	—			

Table S7 Comparison of experimental yields of mono-Boc and mono-Et products with theoretical ones based on ΔG calculations.

	Experimental yield of the reactions ¹			Theoretical, ΔG [kcal/mol]		
	U	T	6-MeU	U	T	6-MeU
	N^1 (3a)	N^1 (2a)	N^1 (1a)	N^1	N^1	N^1
mono-Boc						

mono-Et	—	33 % 1.5eq D	5 % 1.5eq D	—11.0	—10.9	2.1
	12 % 3.0eq D	59 % 3.0eq D	3 % 3.0eq D			
	—	—	9 % 3.0eq P			
	N^3 (3b)	N^3 (2b)	N^3 (1b)	N^3	N^3	N^3
	—	4 % 3.0eq P	5 % 1.5eq D	—11.0	—10.1	—10.5
	12 % 3.0eq D	—	3 % 3.0eq D			
	23 % 3.0eq P	—	15 % 3.0eq P			
	N^1 (3h)	N^1 (2h)	N^1	N^1	N^1	N^1
	—	19 % 1.0eq	—	—39.0	—28.0	—23.7
	12 % 1,5 eq	22 % 1.5eq	—			
	16 % 3,0 eq	33 % 3.0 eq	—			
	N^3 (3i)	N^3	N^3 (1i)	N^3	N^3	N^3
	—	—	17 % 1.0eq	—38.6	—27.6	—27.2
	1 % 1.5eq	—	5 % 1.5eq			
	1 % 3.0eq	—	—			

Table S8 The molecular structure of the **1g** product for the reaction of pyridine (**4**) and Boc₂O with 3-Boc-6-methyluracil (**1b**) according to the B3LYP/6-31G(d,p) calculations. Geometry readable by the JMol software.

C	1.916592	4.482851	0.867407
C	2.1090113	0.49713	0.453690
C	1.098484	2.177506	0.163332
C	1.430283	0.814915	-0.223707
N	2.817325	0.515411	-0.259345
C	3.854049	1.382709	0.36960
N	3.424213	2.643926	0.381503
H	4.158033	3.305004	0.602001
O	5.028663	1.059453	-0.002887
C	3.190759	-0.836639	-0.677551
O	3.403014	-1.102829	-1.826657
O	3.235844	-1.612180	0.382780
C	3.573083	-3.073560	0.281969
C	2.523637	-3.772164	-0.576935
H	2.687517	-4.851820	-0.534051
H	1.518519	-3.566617	-0.200493
H	2.577864	-3.460616	-1.619517
C	4.990615	-3.224458	-0.261231
H	5.289474	-4.273720	-0.196132
H	5.062336	-2.913831	-1.303090
H	5.694761	-2.634564	0.331075
C	3.486398	-3.522896	1.735761
H	3.708901	-4.589926	1.805864
H	4.204598	-2.980985	2.355704
H	2.484331	-3.354644	2.137266
O	0.640658	-0.070817	-0.515087
C	-0.373713	2.582748	0.234069
C	-1.076924	2.486177	-1.103463
C	-2.162030	1.741981	-1.317759
N	-2.768708	0.974909	-0.303285
C	-3.882537	0.202925	-0.626181
O	-4.331432	-0.461996	0.444265
C	-5.523353	-1.350930	0.373016

C	-5.257565	-2.498327	-0.599357
H	-5.185497	-2.146288	-1.627342
H	-4.330577	-3.016567	-0.339165
H	-6.075294	-3.221329	-0.538771
C	-5.631120	-1.866439	1.805189
H	-4.729015	-2.411742	2.093146
H	-5.775548	-1.041278	2.506940
H	-6.483860	-2.543529	1.893083
C	-6.756150	-0.531731	-0.003177
H	-6.694590	-0.156240	-1.023699
H	-7.647603	-1.158961	0.080039
H	-6.876634	0.315775	0.677048
O	-4.350846	0.165633	-1.747268
C	-2.230695	1.066863	0.996942
C	-1.148237	1.796805	1.270686
H	-0.800719	1.833206	2.297999
H	-2.767054	0.516592	1.753976
H	-2.657314	1.681083	-2.275649
H	-0.673230	3.059377	-1.931675
H	-0.378593	3.634815	0.538884
H	2.872507	4.971975	1.059777
H	1.400185	5.048429	0.088536
H	1.315955	4.550299	1.776993

Table S9 ΔG free energy, kcal/mol, of the reactions leading to the formation of the **1g** product according to the B3LYP/6–31G(d) calculations.

Reaction	Substrates	Products	ΔG
R1	4 + (Boc) ₂ O	5 ⁽⁺⁾ + Boc ⁽⁻⁾	21.8
R2	5 ⁽⁺⁾ + Boc ⁽⁻⁾ + 1b	6 ¹	40.7
R3	6	1g + CO ₂ + <i>tert</i> -Butanol	-48.2
		Sum of ΔG	14.3 ¹⁾

¹⁾ The geometry of **6** is given in the Table S7.

Table S10 The molecular structure of the transition complex (**6**) for the reaction of pyridine (**4**) and Boc₂O with 3–Boc–6–methyluracil (**1b**), Scheme 2, according to the B3LYP/6–31G(d) calculations. Geometry readable by the Jmol software.

C	-2.257480	0.519040	3.476540
C	-1.510368	1.182451	2.374032
C	-1.142166	0.462009	1.166957
C	-0.413417	1.261447	0.128385
N	-0.166848	2.620945	0.426448
C	-0.556162	3.284099	1.563966
N	-1.256270	2.477031	2.493697
O	-0.347490	4.456894	1.790249
C	0.501958	3.427685	-0.612803
O	-0.138861	4.034428	-1.430716
O	1.798625	3.330414	-0.439698
C	2.772631	4.027442	-1.354044

C	2.588994	3.496897	-2.773897
C	2.563884	5.535684	-1.241548
C	4.116157	3.603407	-0.767969
O	-0.051380	0.829067	-0.948667
C	-0.501558	-0.979471	1.437985
C	-0.518818	-1.871254	0.225366
C	0.574117	-2.470304	-0.259551
N	1.854312	-2.295264	0.303786
C	2.939359	-2.959692	-0.274852
O	4.073914	-2.679735	0.377708
C	5.377008	-3.263277	-0.035311
C	5.724776	-2.801928	-1.451403
C	6.341172	-2.660202	0.986015
C	5.326178	-4.786062	0.096640
O	2.815046	-3.688055	-1.246603
C	1.962063	-1.463205	1.429204
C	0.899747	-0.858031	1.978828
O	-3.739426	0.069263	0.489936
C	-4.220333	-0.869204	-0.233047
O	-5.597109	-0.757821	-0.308351
C	-6.410709	-1.702109	-1.068767
C	-7.834490	-1.178843	-0.848736
C	-6.054098	-1.647202	-2.559132
C	-6.276047	-3.118332	-0.496229
O	-3.609295	-1.778502	-0.814871
H	-1.575451	2.983576	3.320116
H	3.390105	3.895589	-3.405483
H	2.654108	2.403715	-2.788234
H	1.631127	3.801909	-3.200924
H	3.355715	6.044563	-1.801624
H	1.600129	5.844052	-1.652751
H	2.625232	5.856356	-0.195997
H	4.924839	4.053922	-1.352078
H	4.212157	3.937303	0.270516
H	4.230435	2.514901	-0.800033
H	5.040147	-3.222672	-2.190308
H	5.691270	-1.708604	-1.517635
H	6.742607	-3.125926	-1.695057
H	6.341061	-1.566646	0.923847
H	6.063013	-2.952346	2.004351
H	7.357385	-3.017266	0.789940
H	4.635385	-5.228454	-0.623659
H	6.326273	-5.195965	-0.082565
H	5.017359	-5.073230	1.108074
H	1.065557	-0.240512	2.857885
H	2.959649	-1.361043	1.831462
H	0.556542	-3.130721	-1.115871
H	-1.477309	-2.035008	-0.261491
H	-1.167527	-1.411560	2.196452
H	-2.677182	1.243139	4.180403
H	-3.059222	-0.088470	3.045220
H	-1.584132	-0.155094	4.021192
H	-2.234622	0.205759	0.724309
H	-8.559834	-1.808182	-1.376435
H	-8.087487	-1.181141	0.217662

H	-7.932212	-0.153039	-1.222006
H	-6.747161	-2.271538	-3.135609
H	-6.135737	-0.618625	-2.931077
H	-5.035451	-2.000217	-2.729040
H	-6.977651	-3.795629	-0.997652
H	-5.261386	-3.497501	-0.629879
H	-6.512673	-3.118267	0.574800

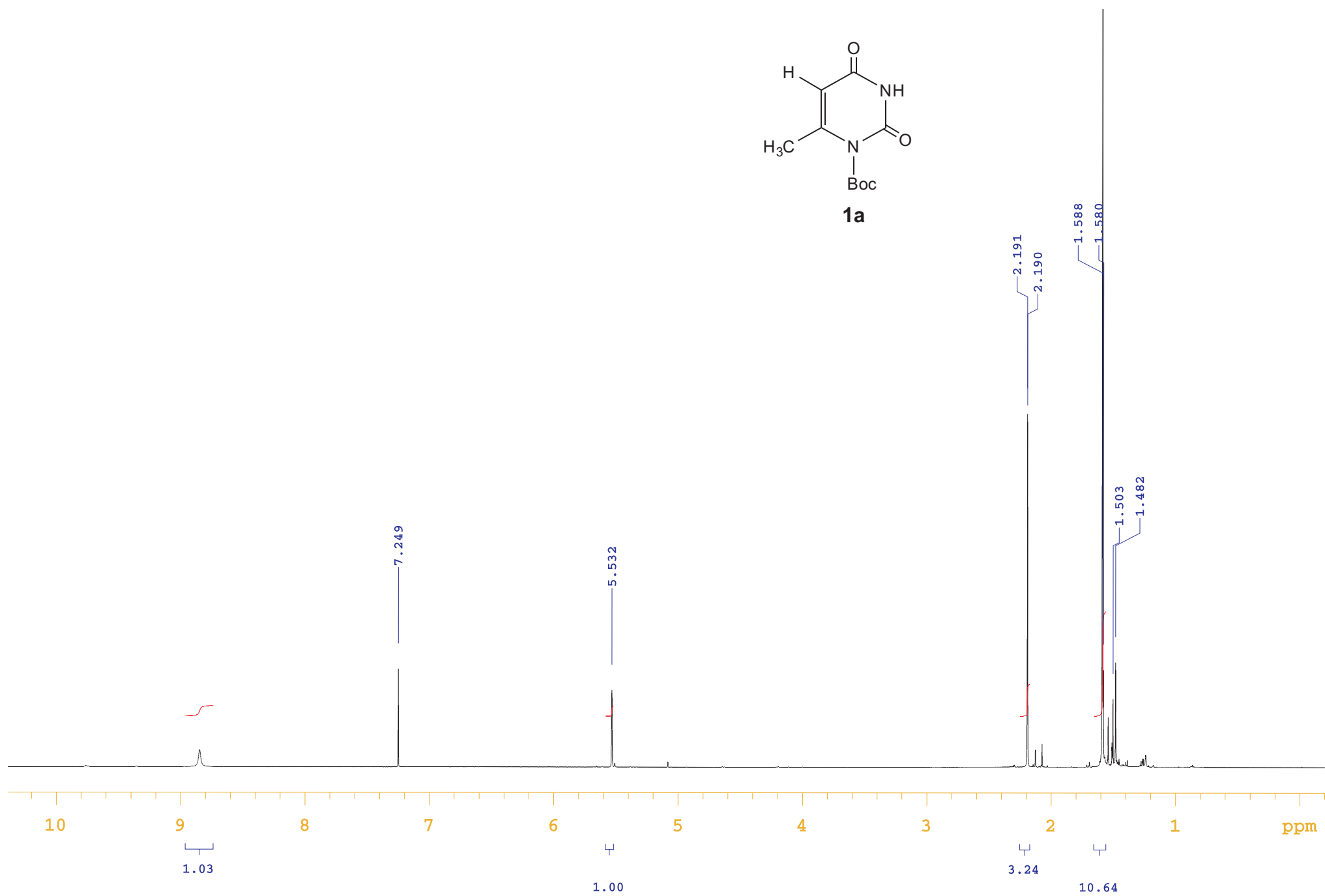
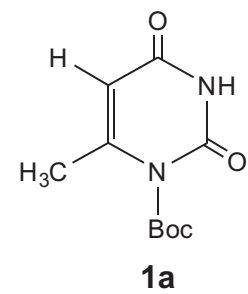
Table S11 The Fukui's condensed function, f , calculated with the DFT/B3LYP/6-31G(d,p) method and using the Mulliken net charges.

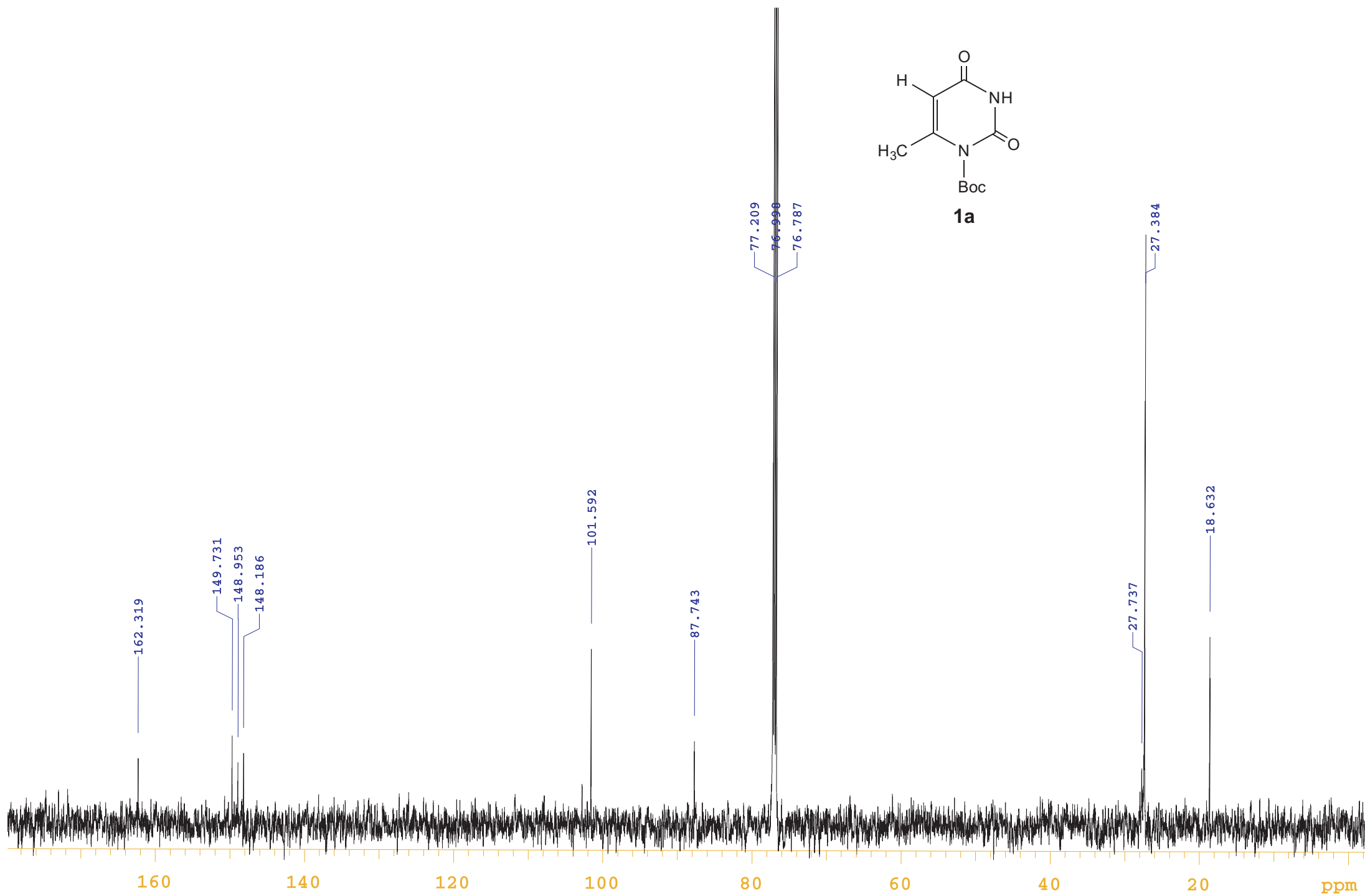
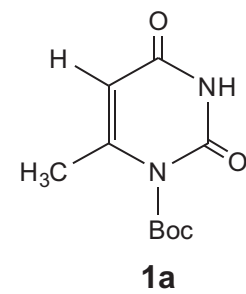
Position in the molecule	Uracil (3)	Thymine (2)	6-methyluracil (1)	<i>N</i> ³ -Boc-6-methyluracil (1b)	<i>N</i> ¹ -Boc-6-methyluracil (1a)
<i>N</i> 1	-0.092	-0.077	-0.081	-0.078	0.050
<i>N</i> 3	0.000	0.003	0.001	0.011	-0.002
<i>O</i> 2	-0.098	-0.096	-0.089	-0.089	-0.080
<i>O</i> 4	-0.106	-0.089	-0.106	-0.104	-0.103
<i>C</i> 5	-0.165	-0.097	-0.186	-0.187	-0.173
<i>C</i> 6	-0.111	-0.133	-0.069	-0.069	-0.067

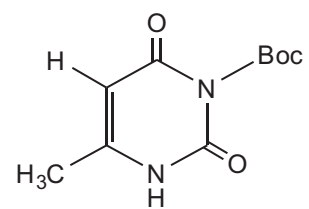
Table S12 The effect of H-bonded pyridine (**4**) and DMAP on the predicted ΔG , in kcal/mol, of reaction of (Boc)₂O with 6-methyluracil.

Product	No explicit solvent B3LYP/6-31G(d,p)	H-bonded Pyridine B3LYP/6-31G(d,p)	H-bonded DMAP B3LYP/6-31G(d,p)
<i>N</i> ¹ -Boc 3a	5.1	5.2	2.3
<i>N</i> ³ -Boc 3b	-7.1	-7.2	-10.9
<i>N</i> ¹ , <i>N</i> ³ -di-Boc 3c	-12.1	-11.7	-21.2
<i>O</i> ² , <i>O</i> ⁴ -di-Boc 3d	-2.7	-2.3	-11.8
<i>O</i> ² , <i>N</i> ³ -di-Boc 3e	2.4	2.8	-6.7

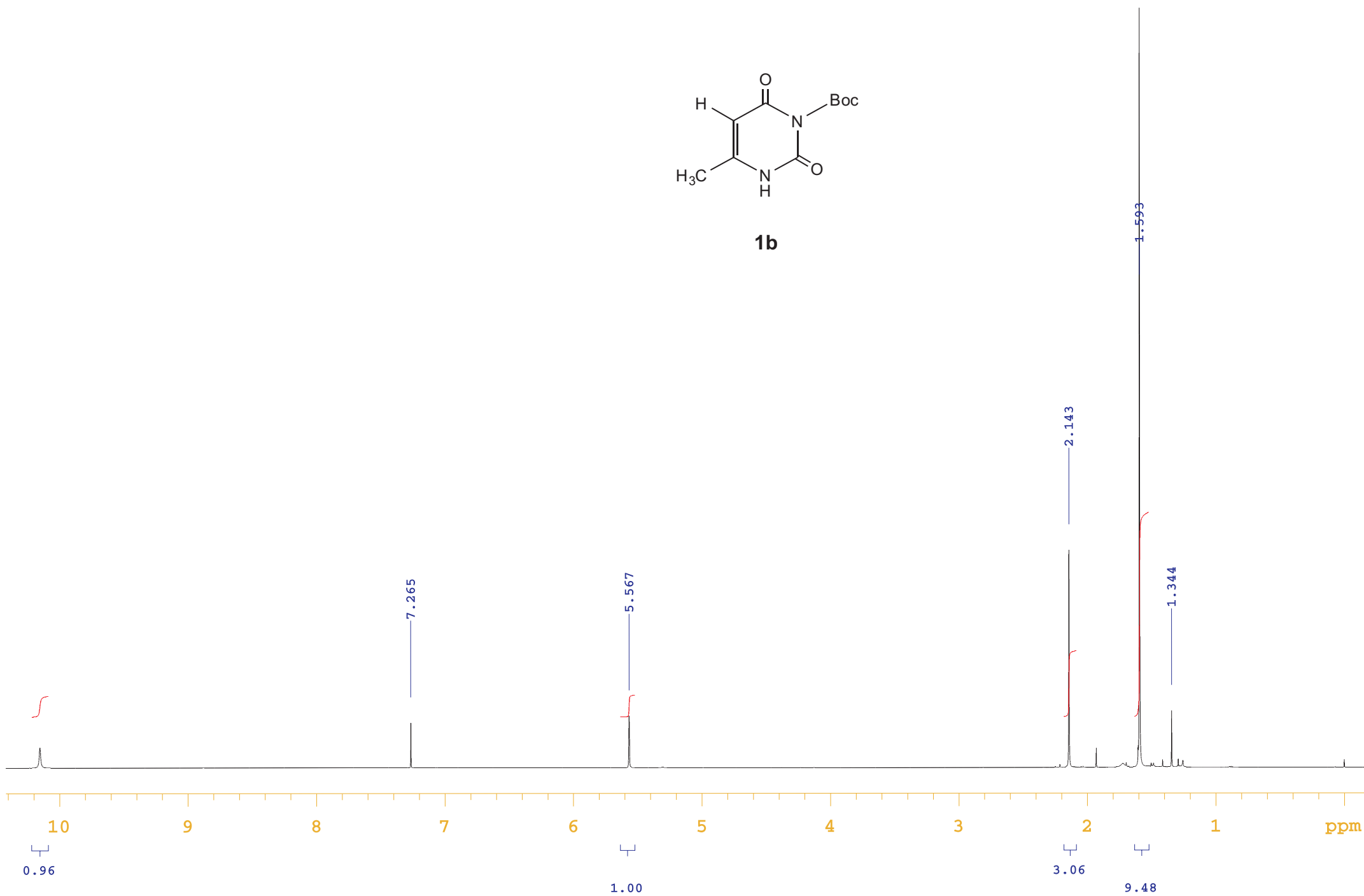
Copies of the ^1H and ^{13}C NMR spectra

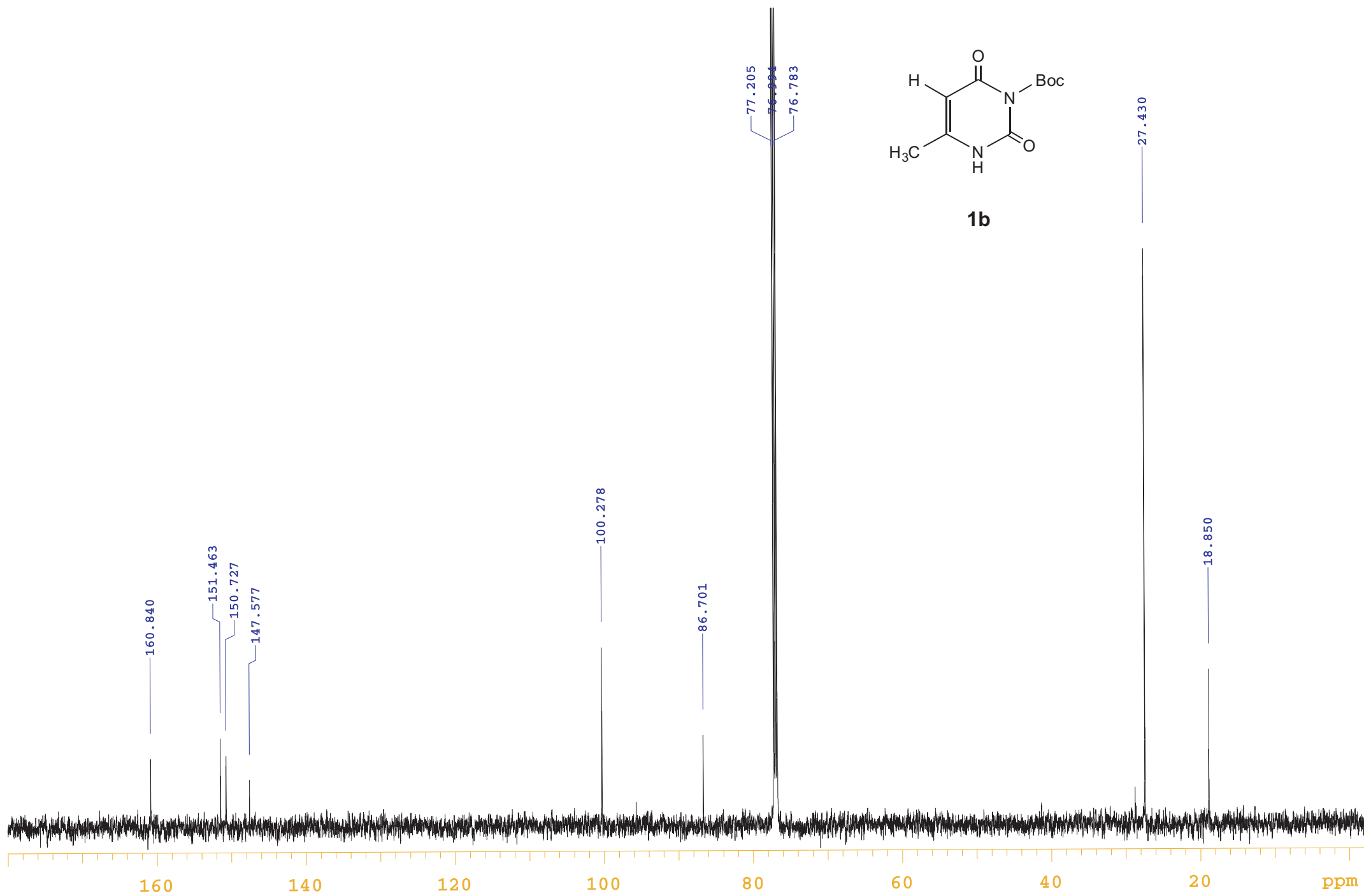


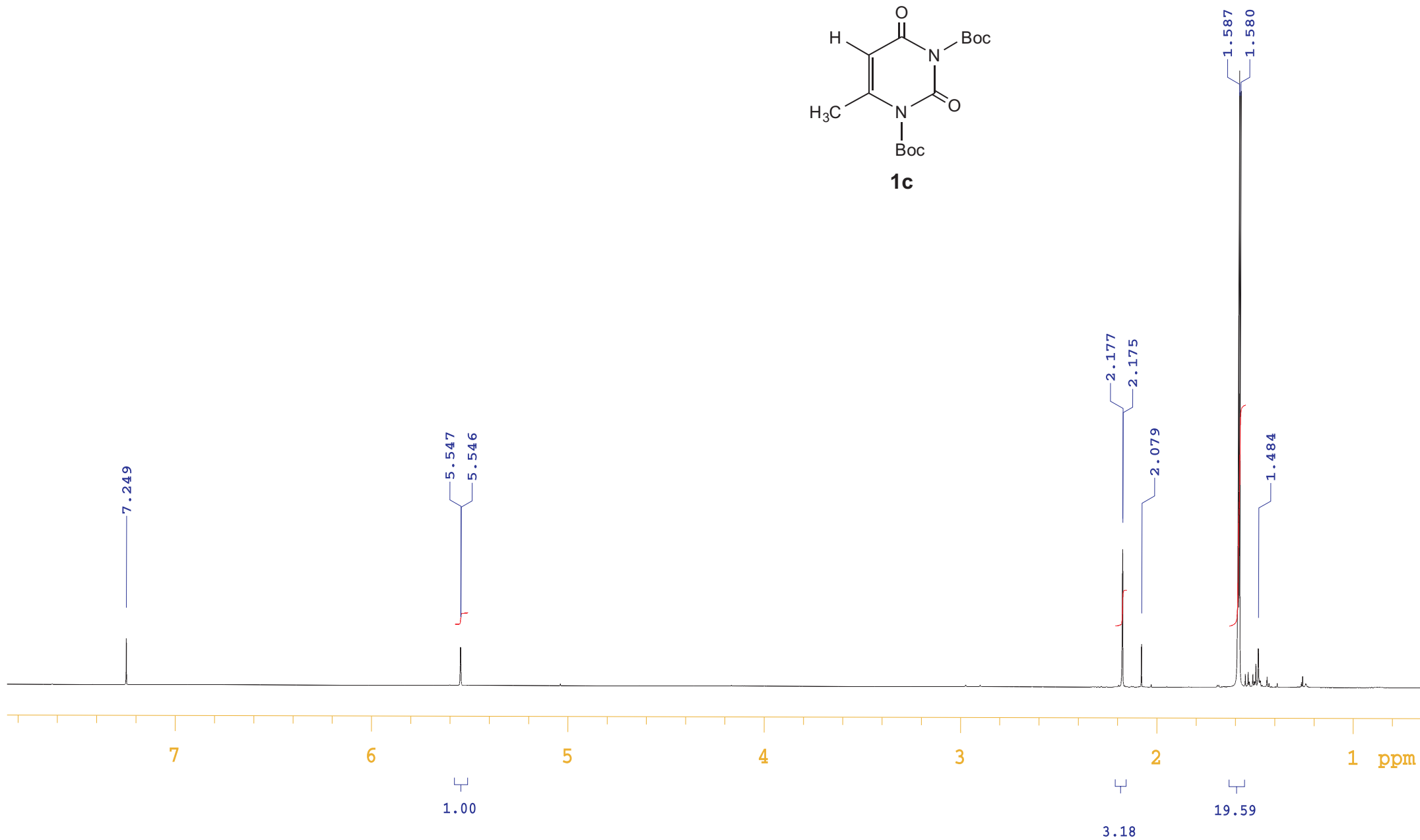
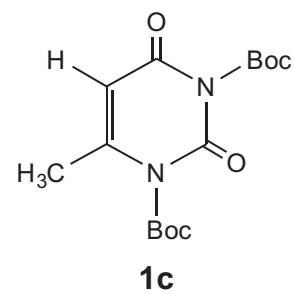


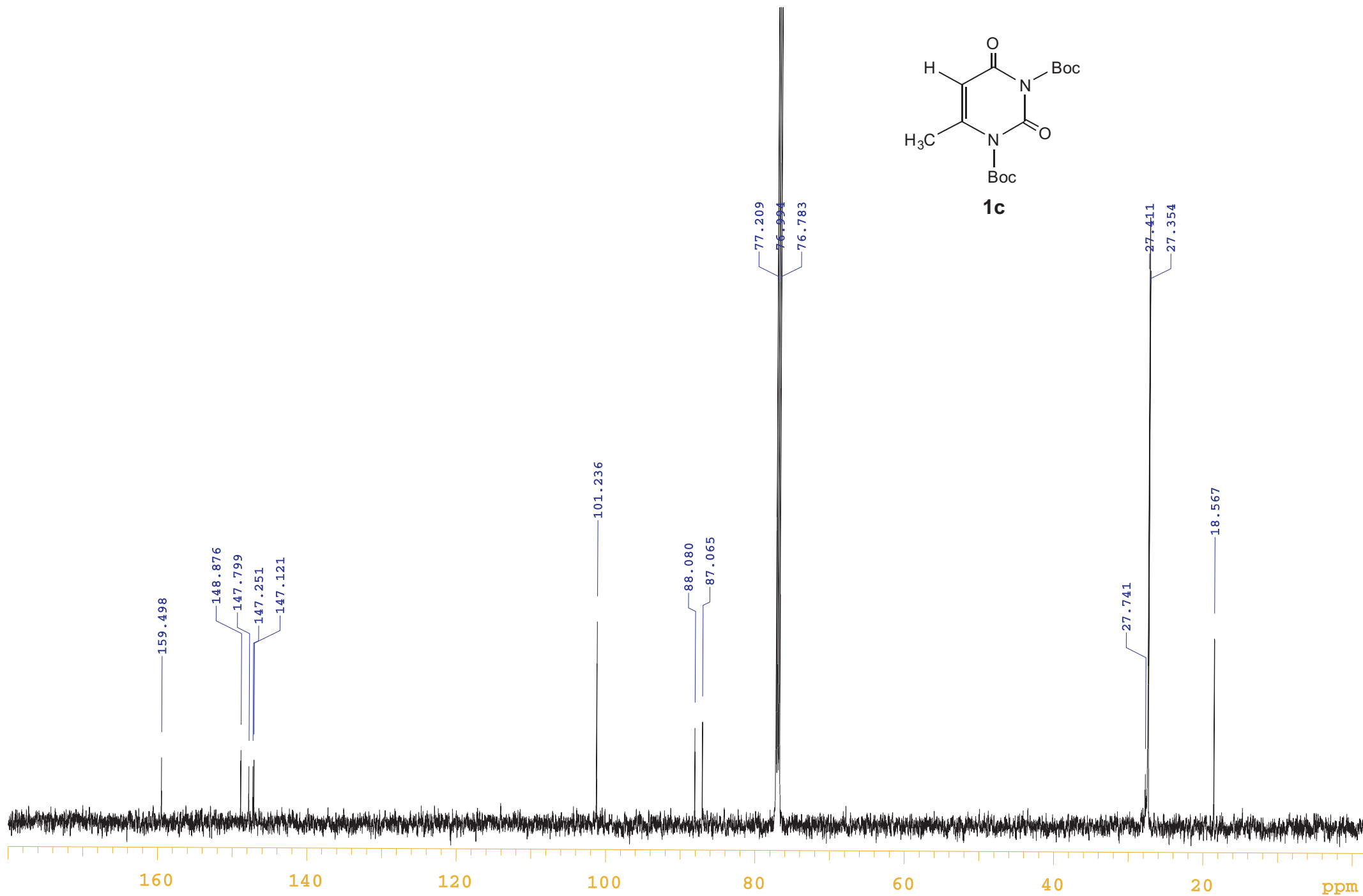
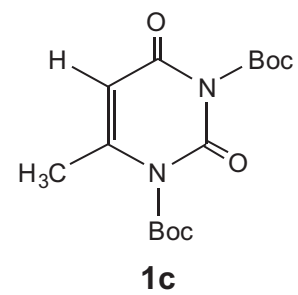


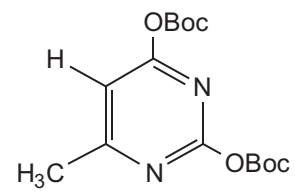
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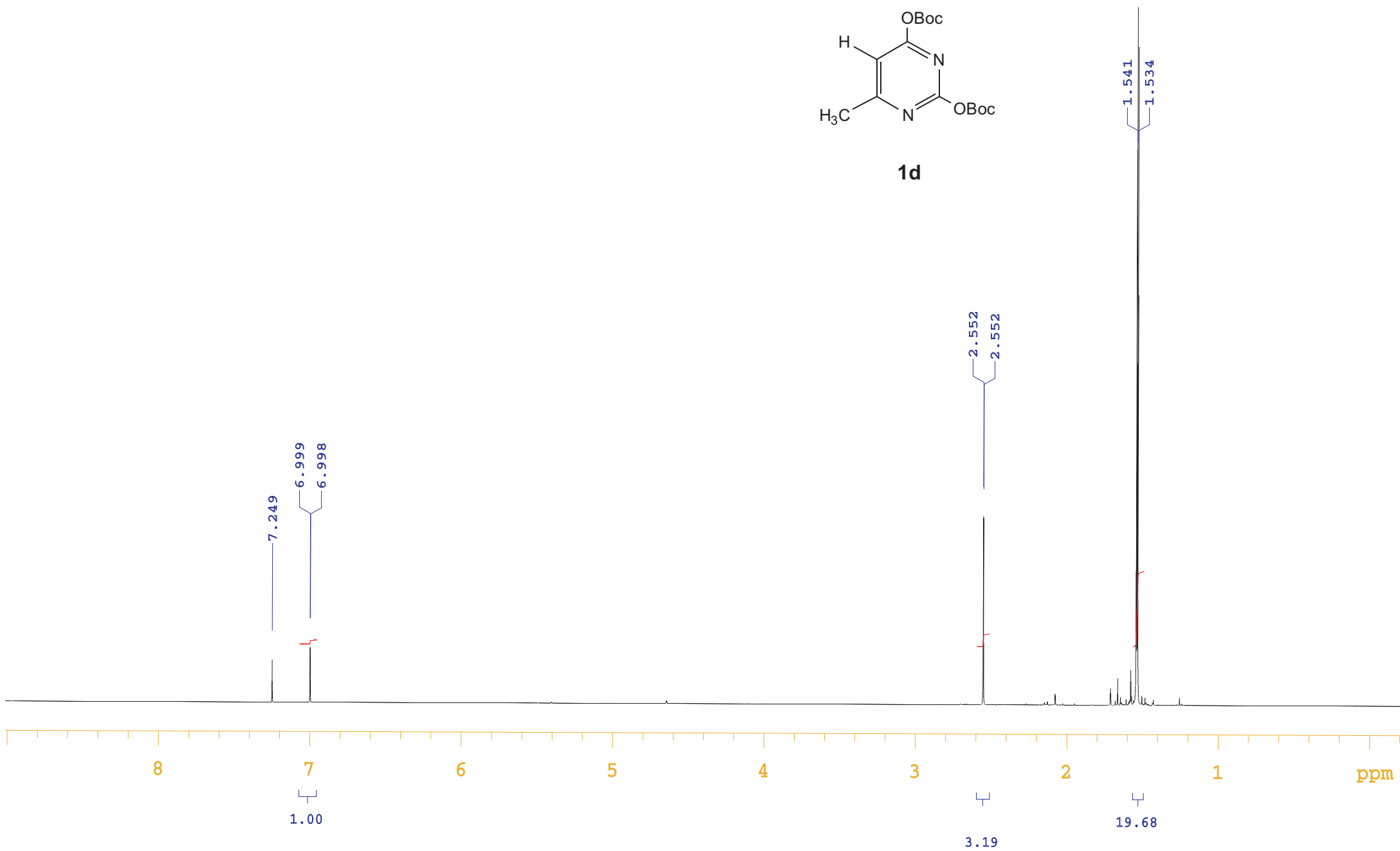


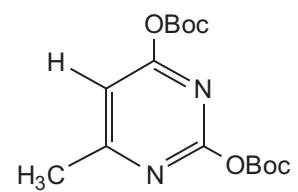




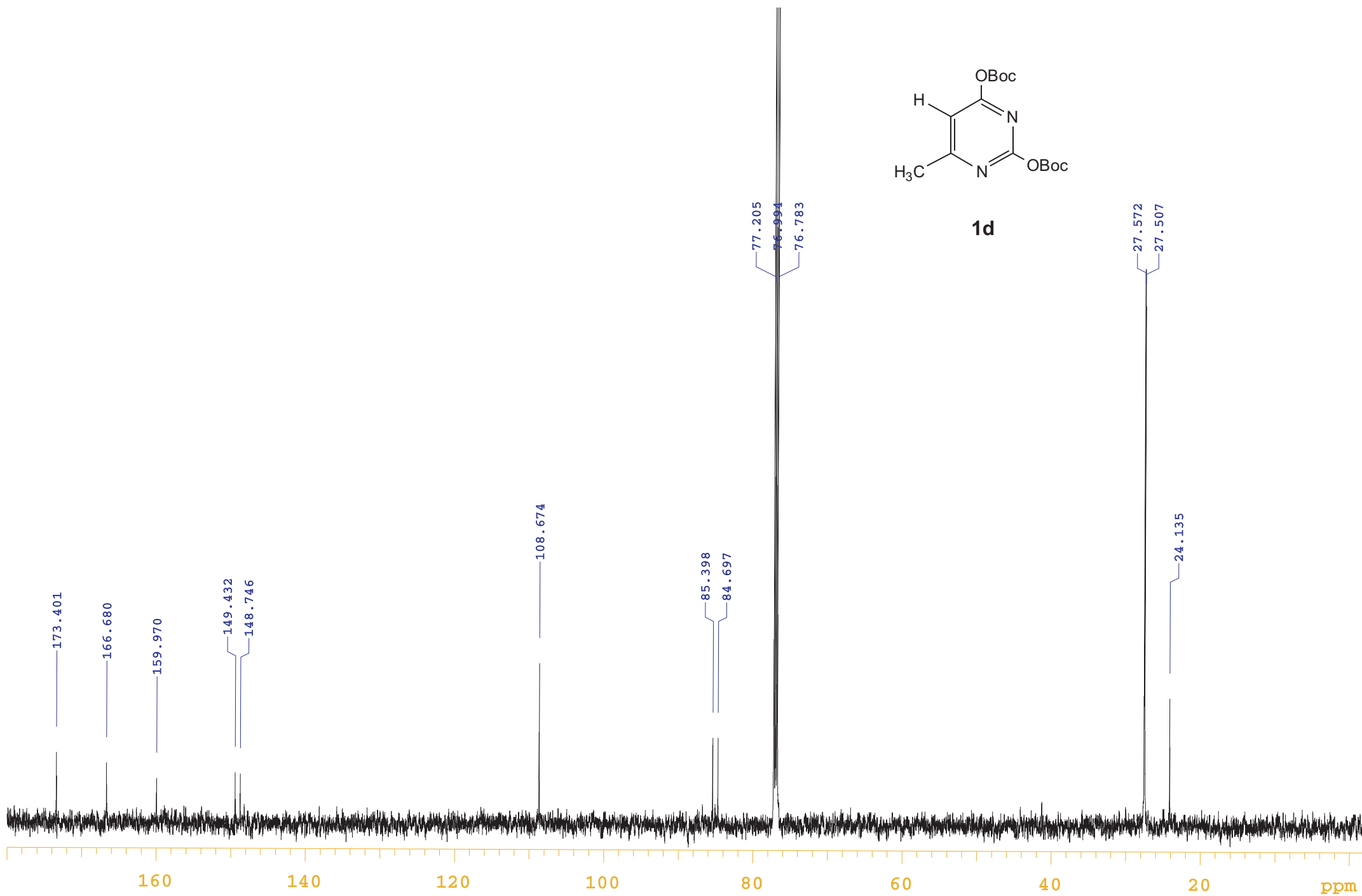


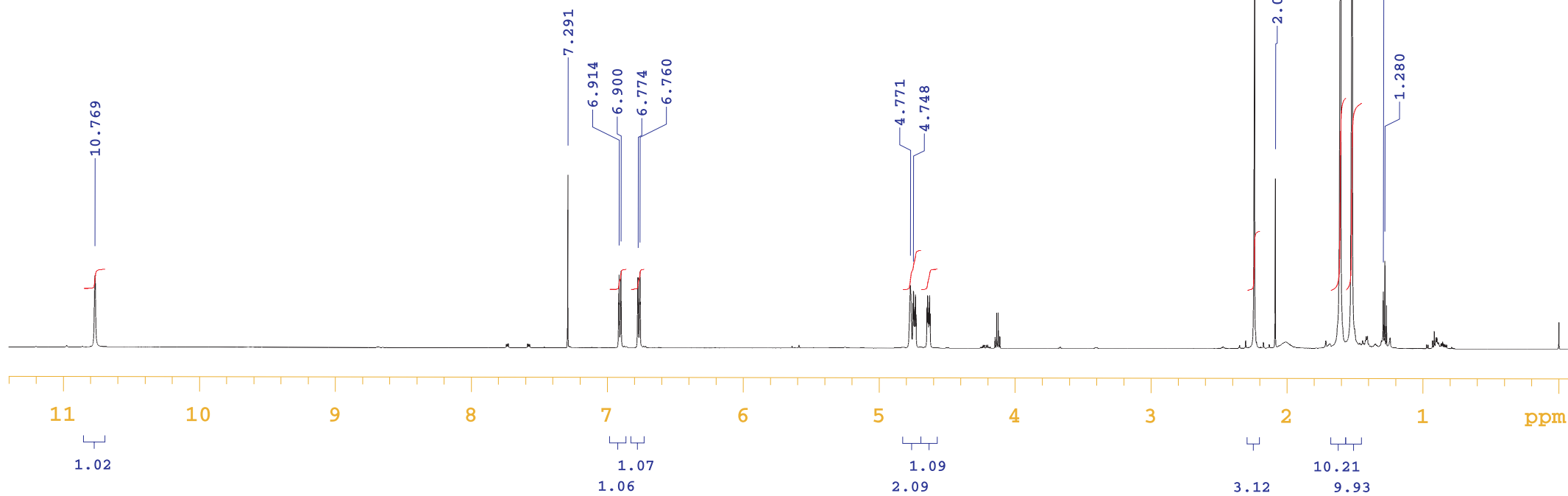
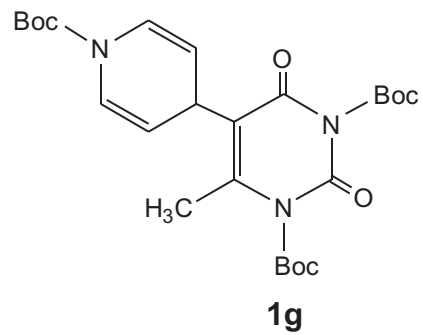
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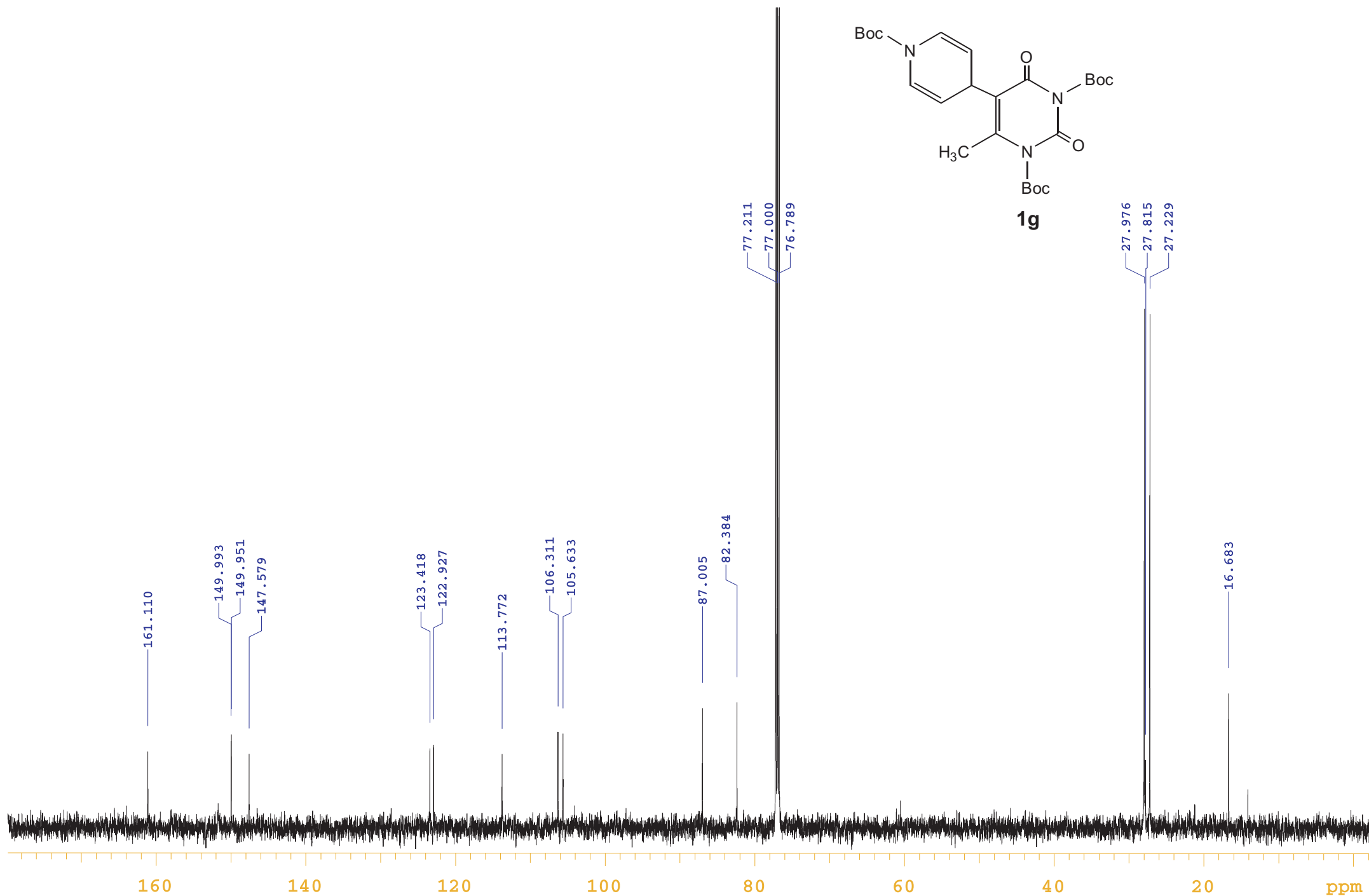
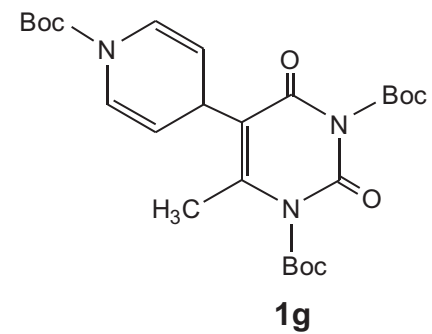


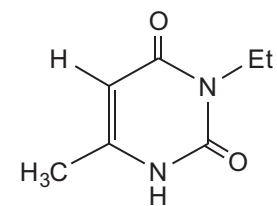


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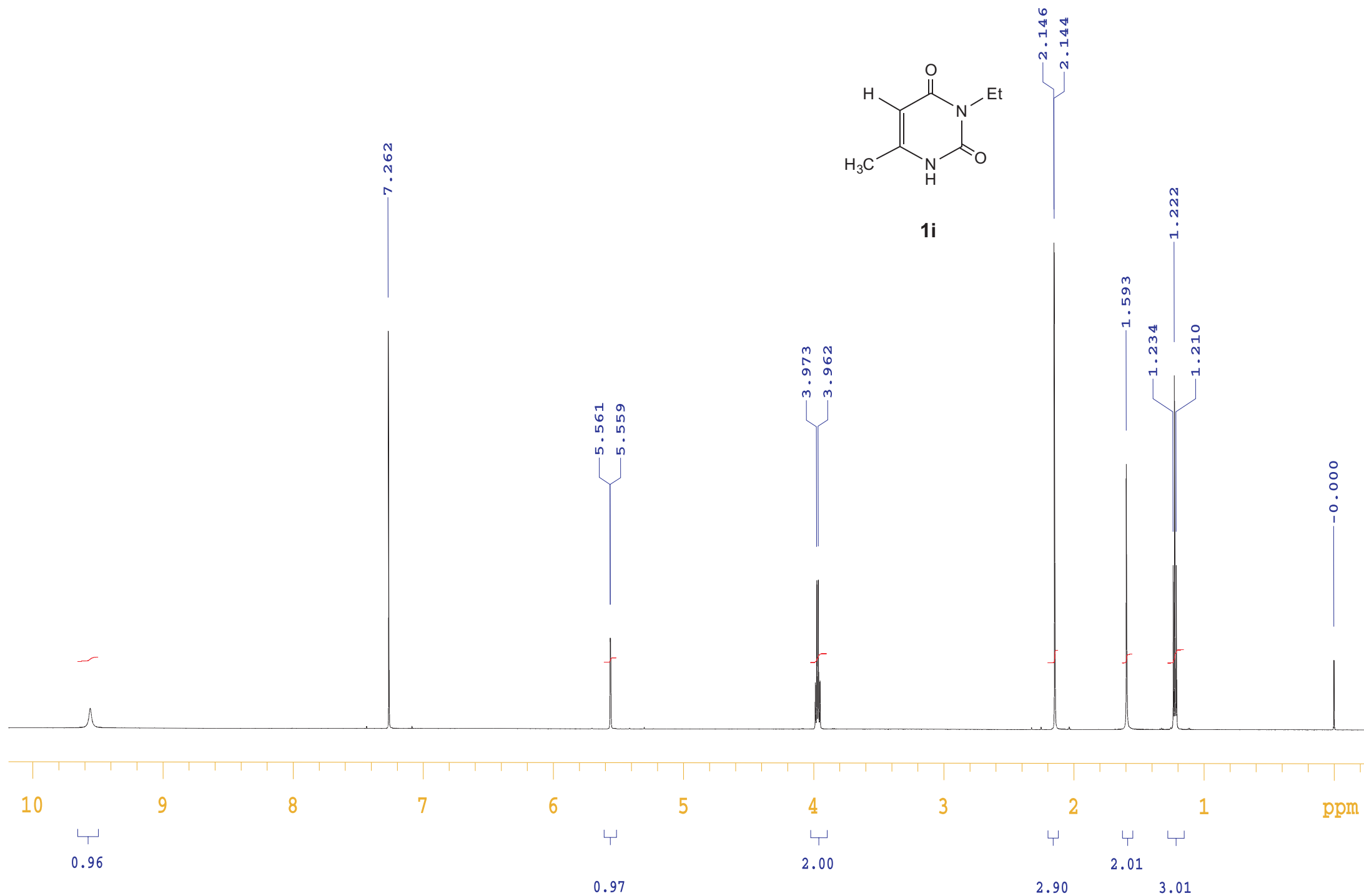


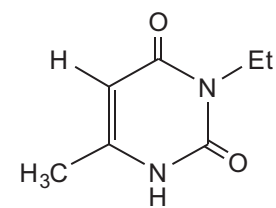




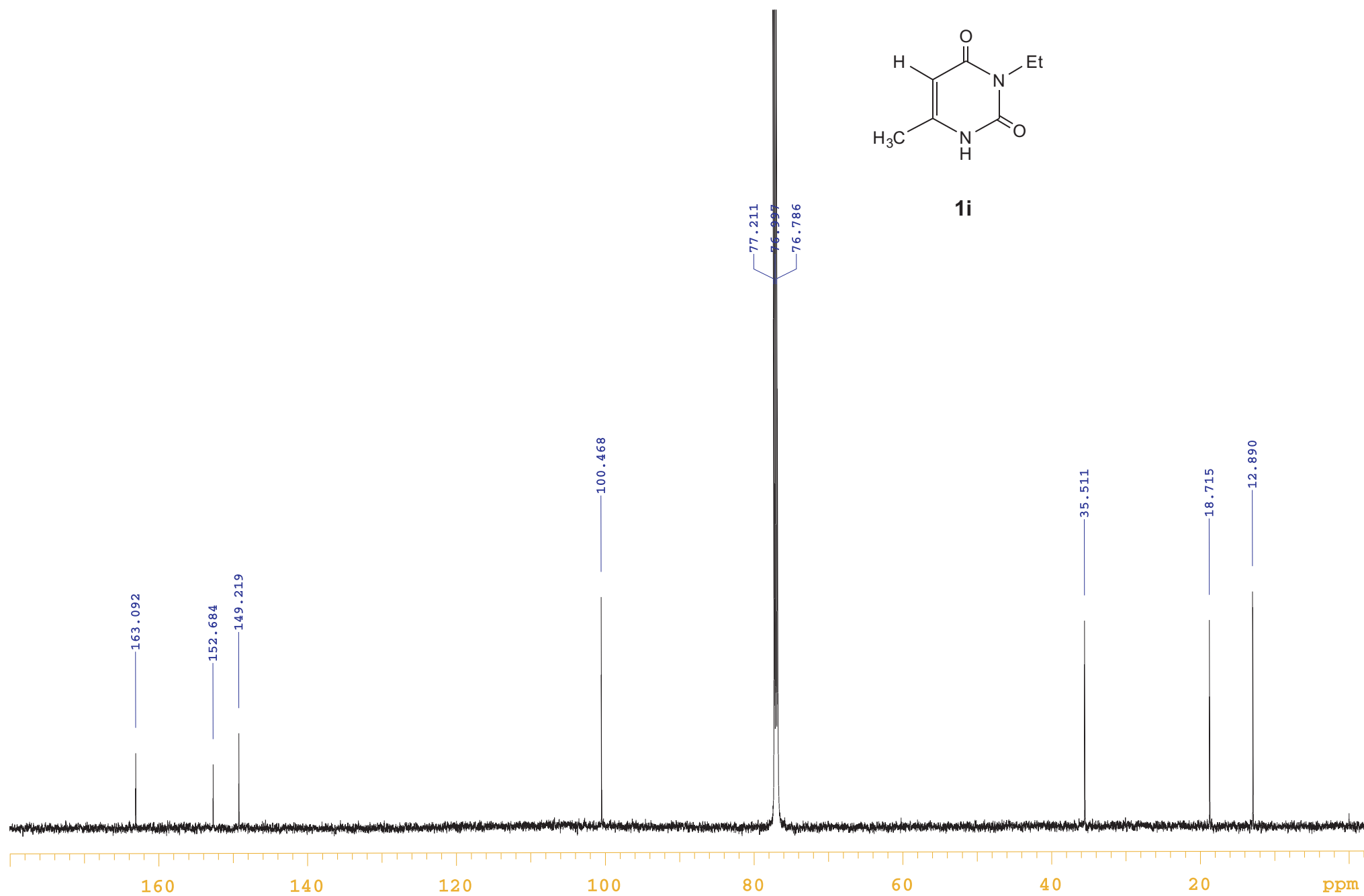


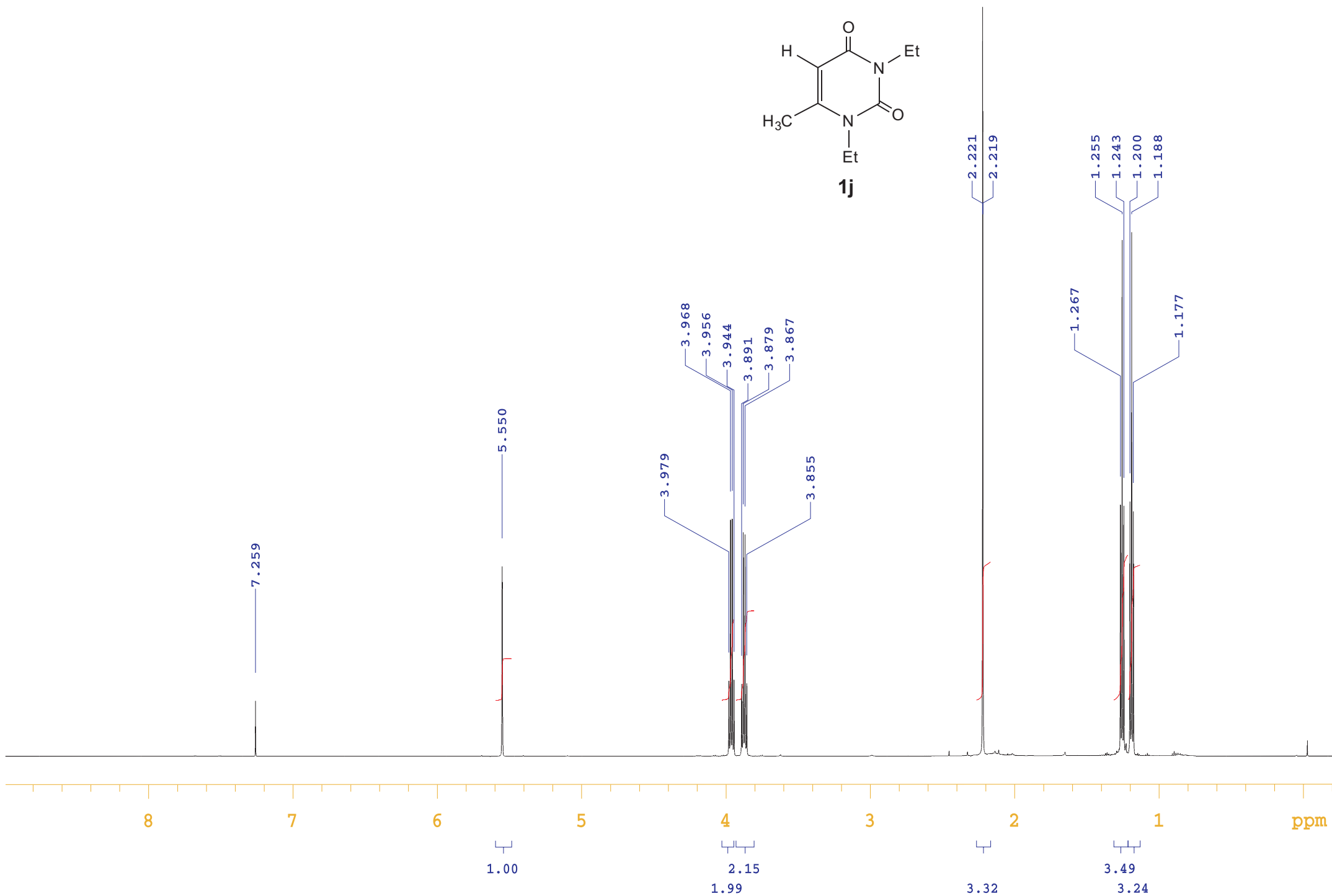
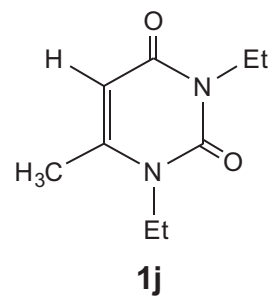
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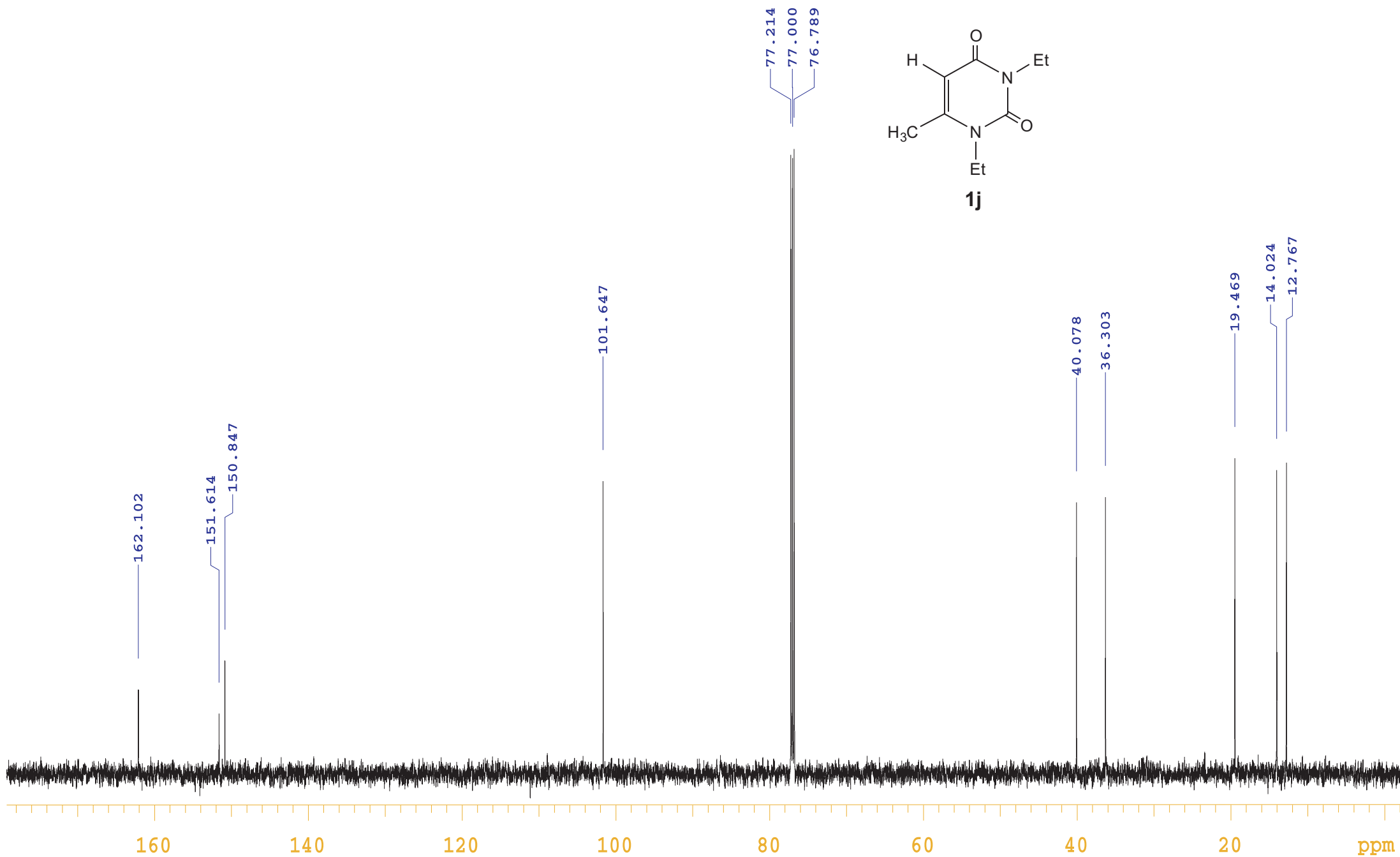


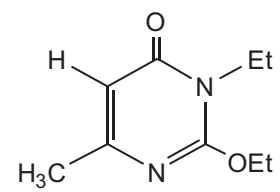


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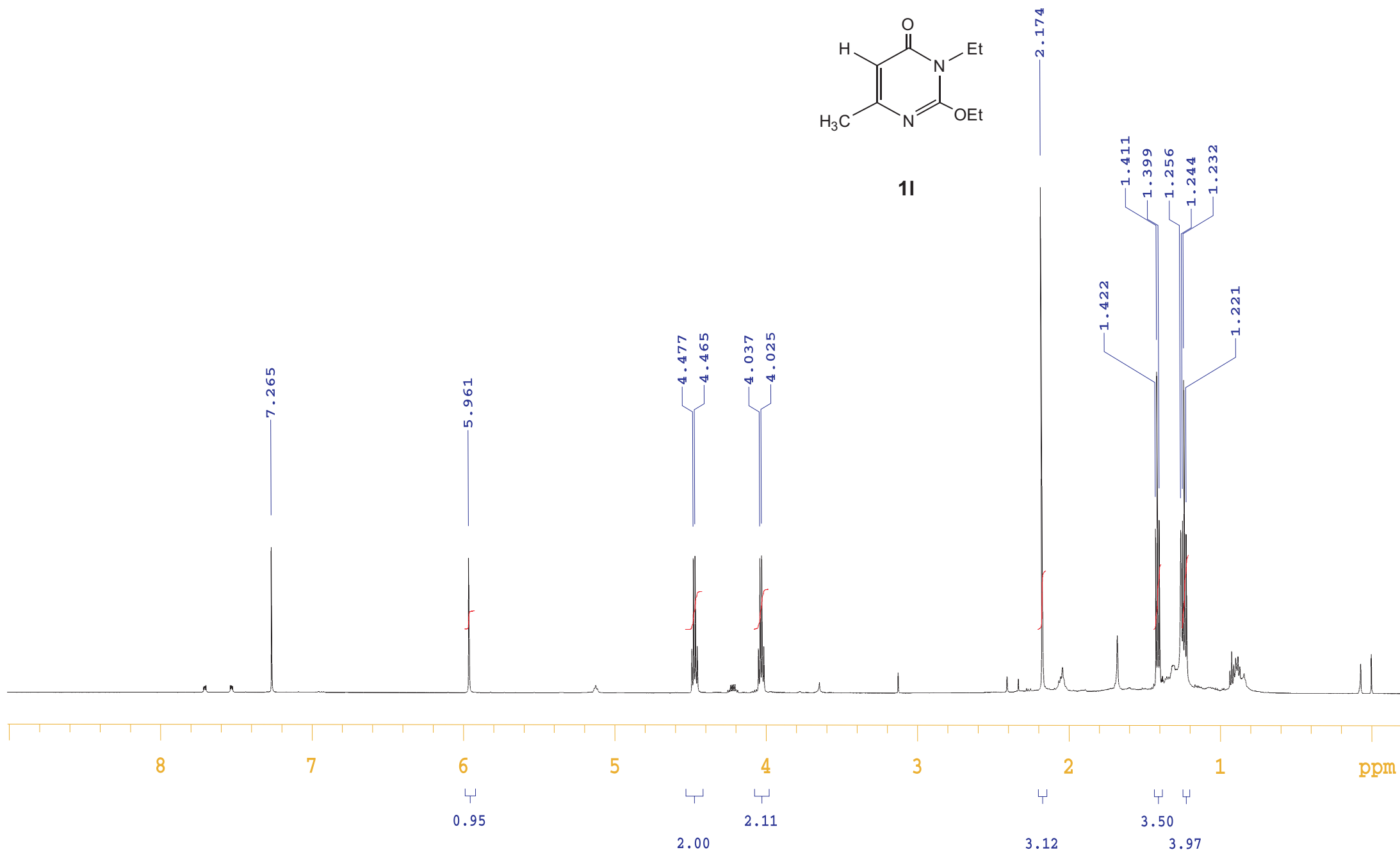


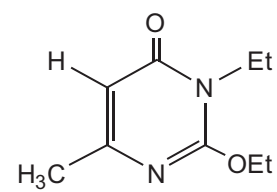




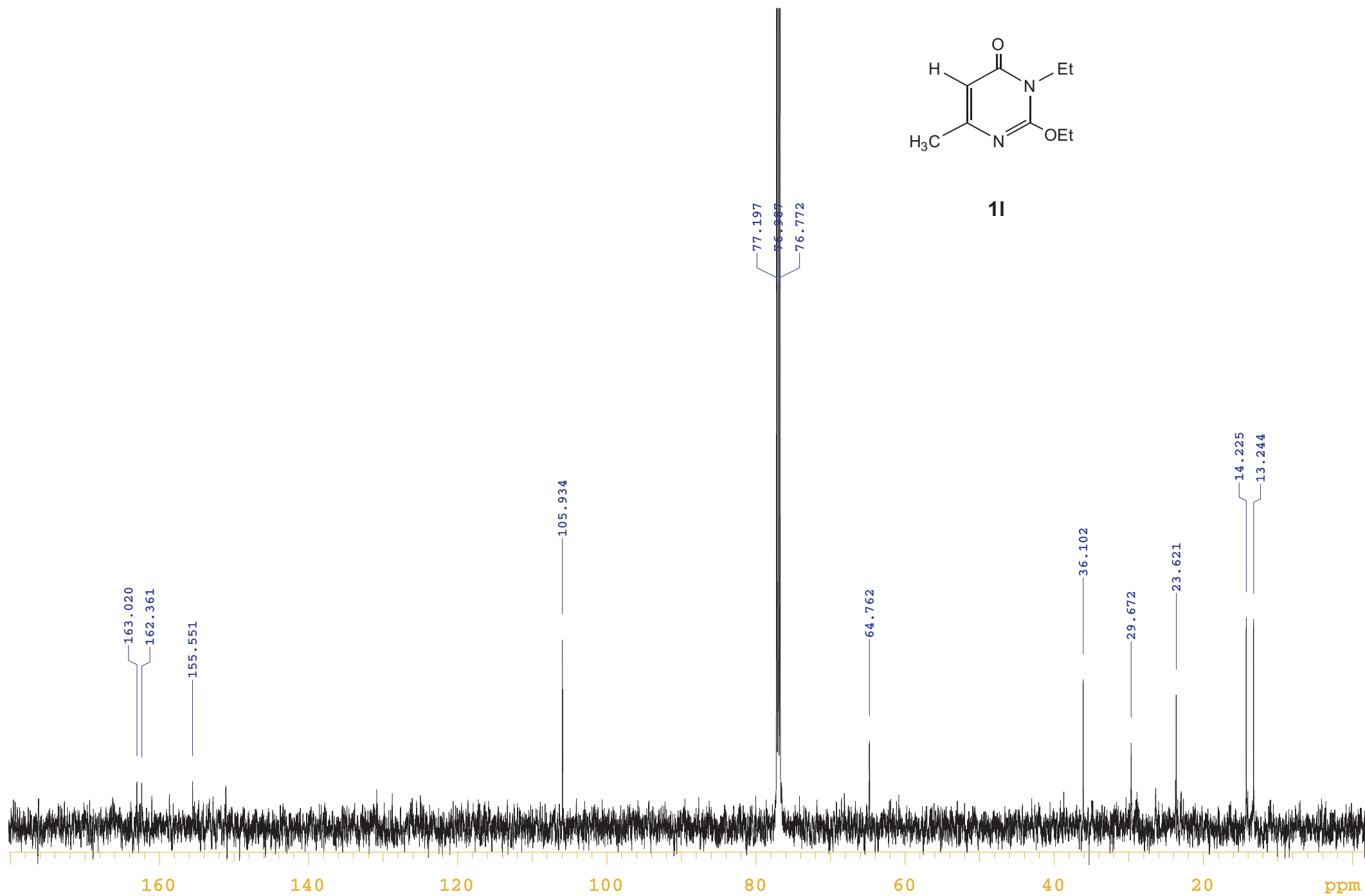


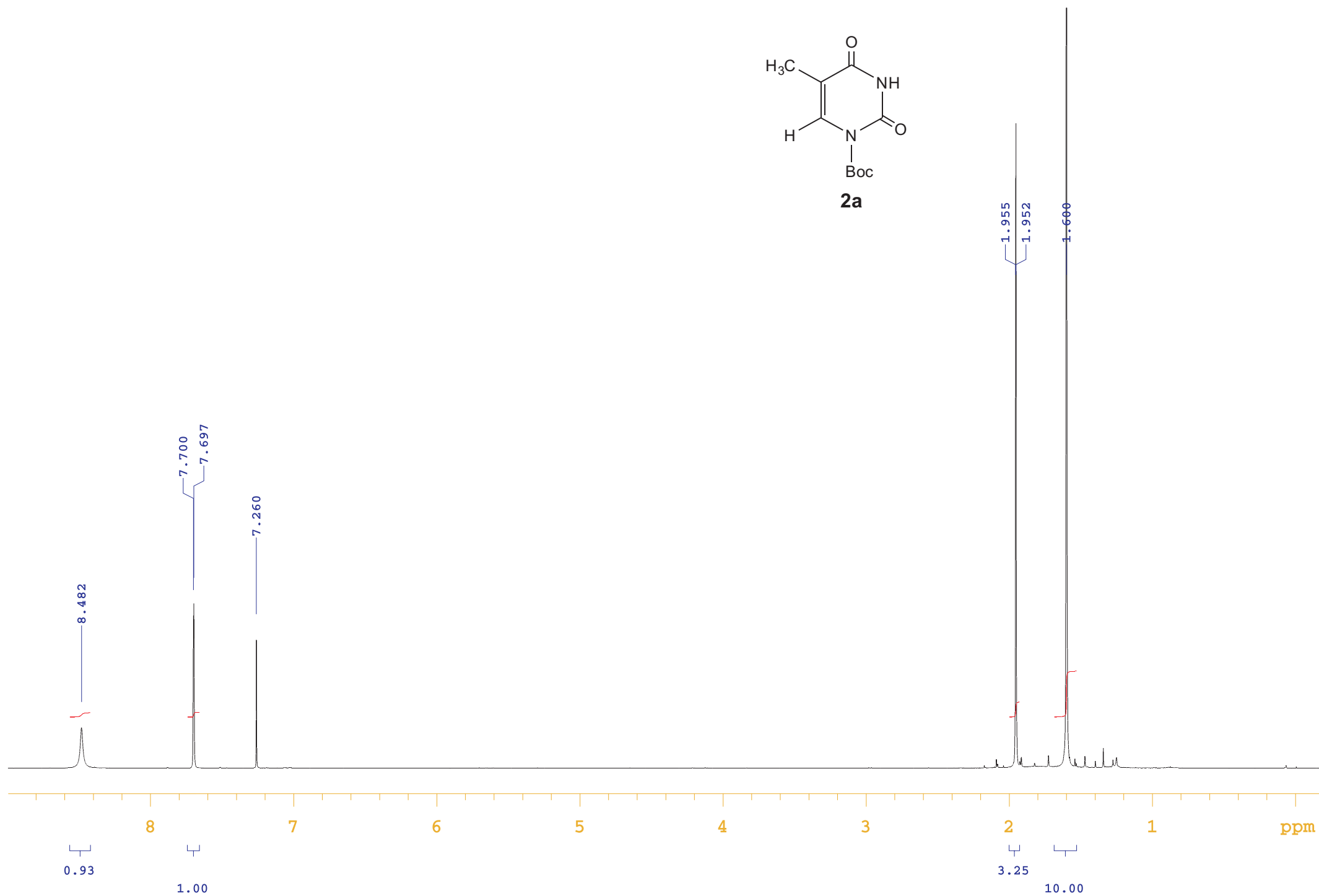
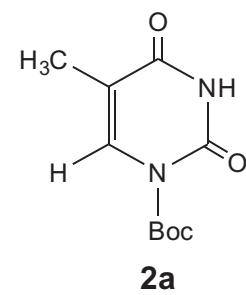
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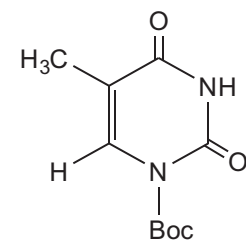




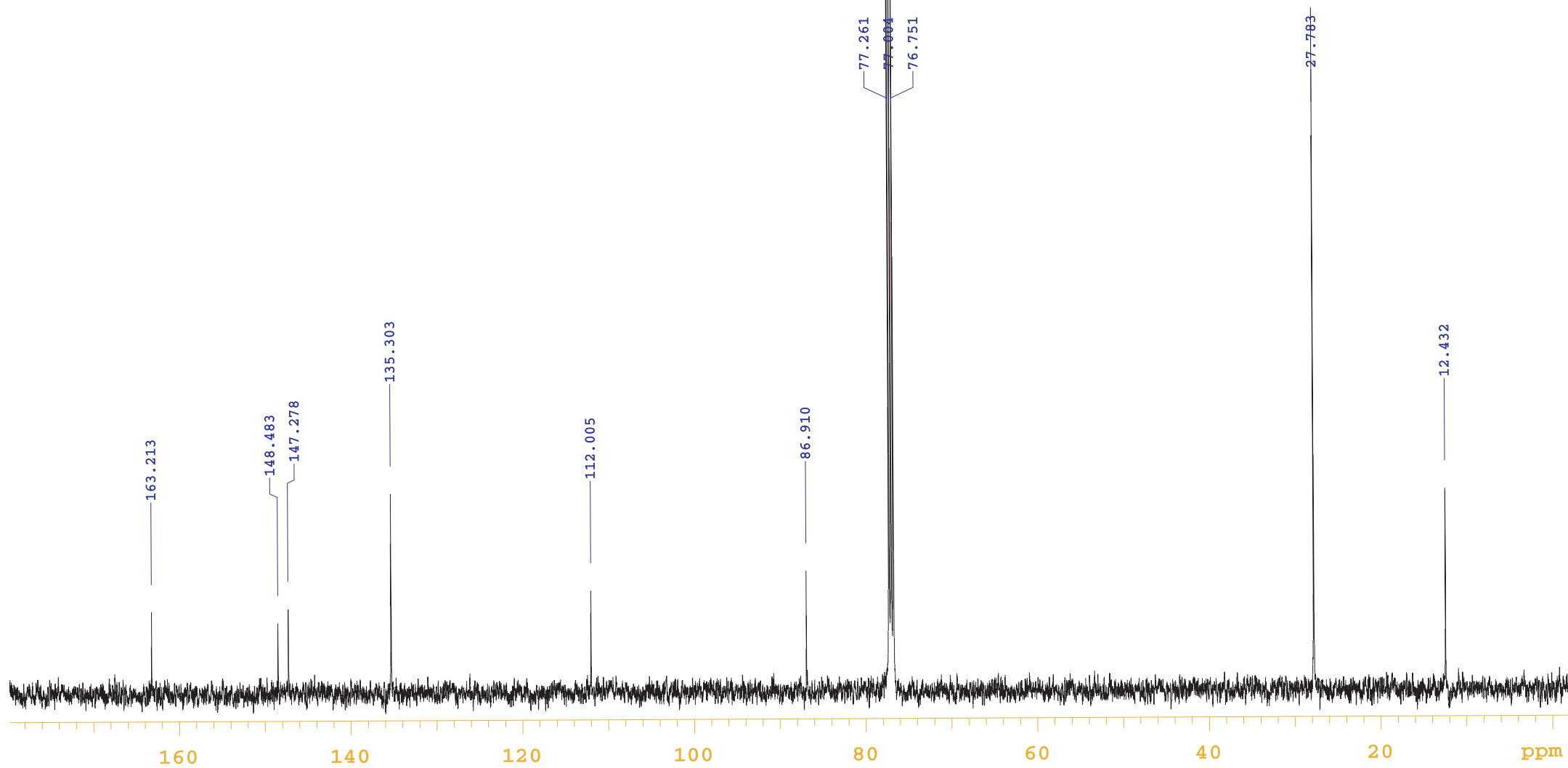
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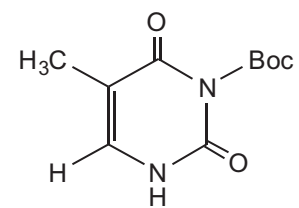




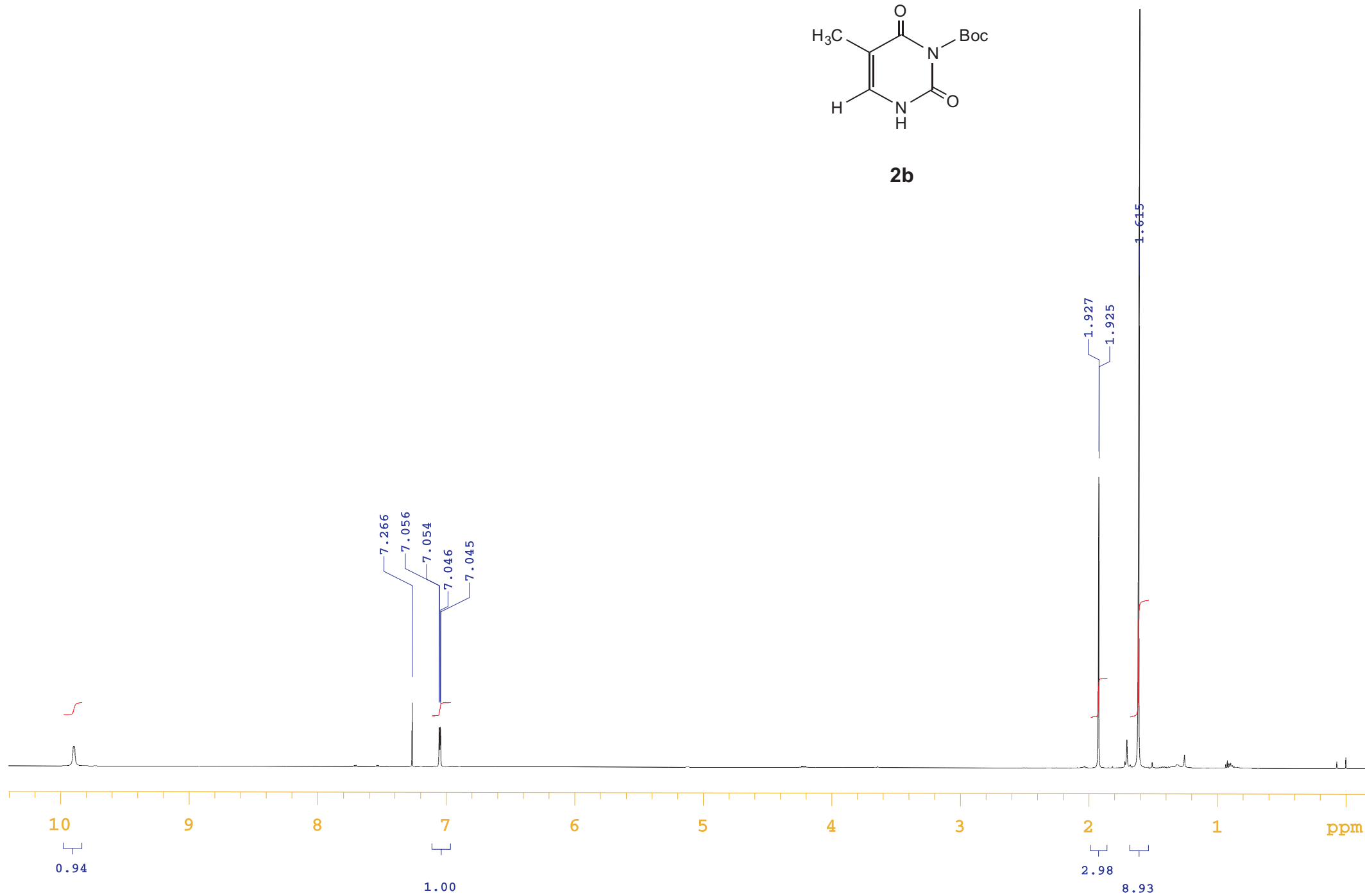


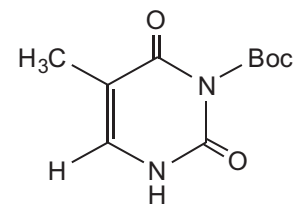
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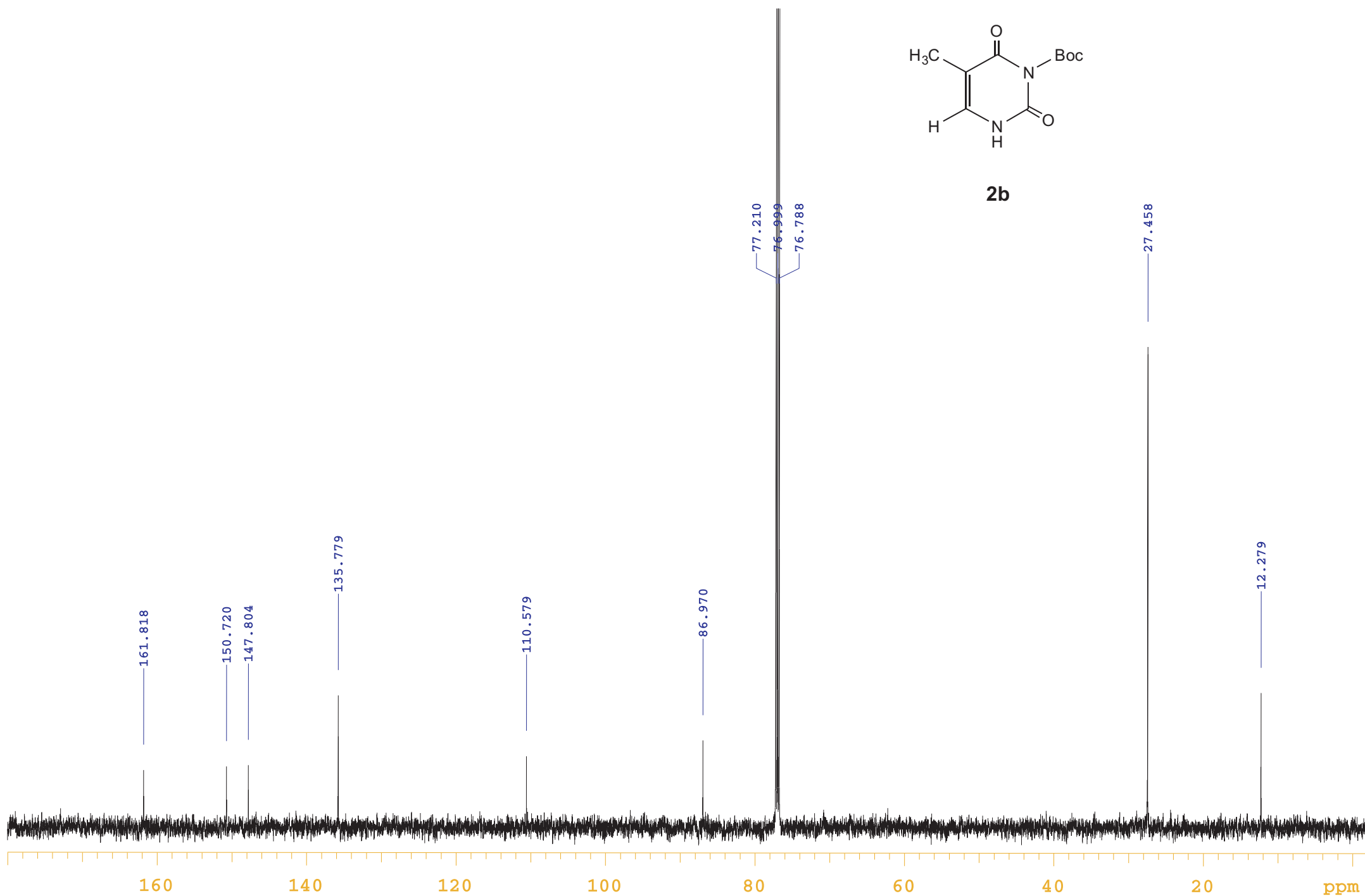


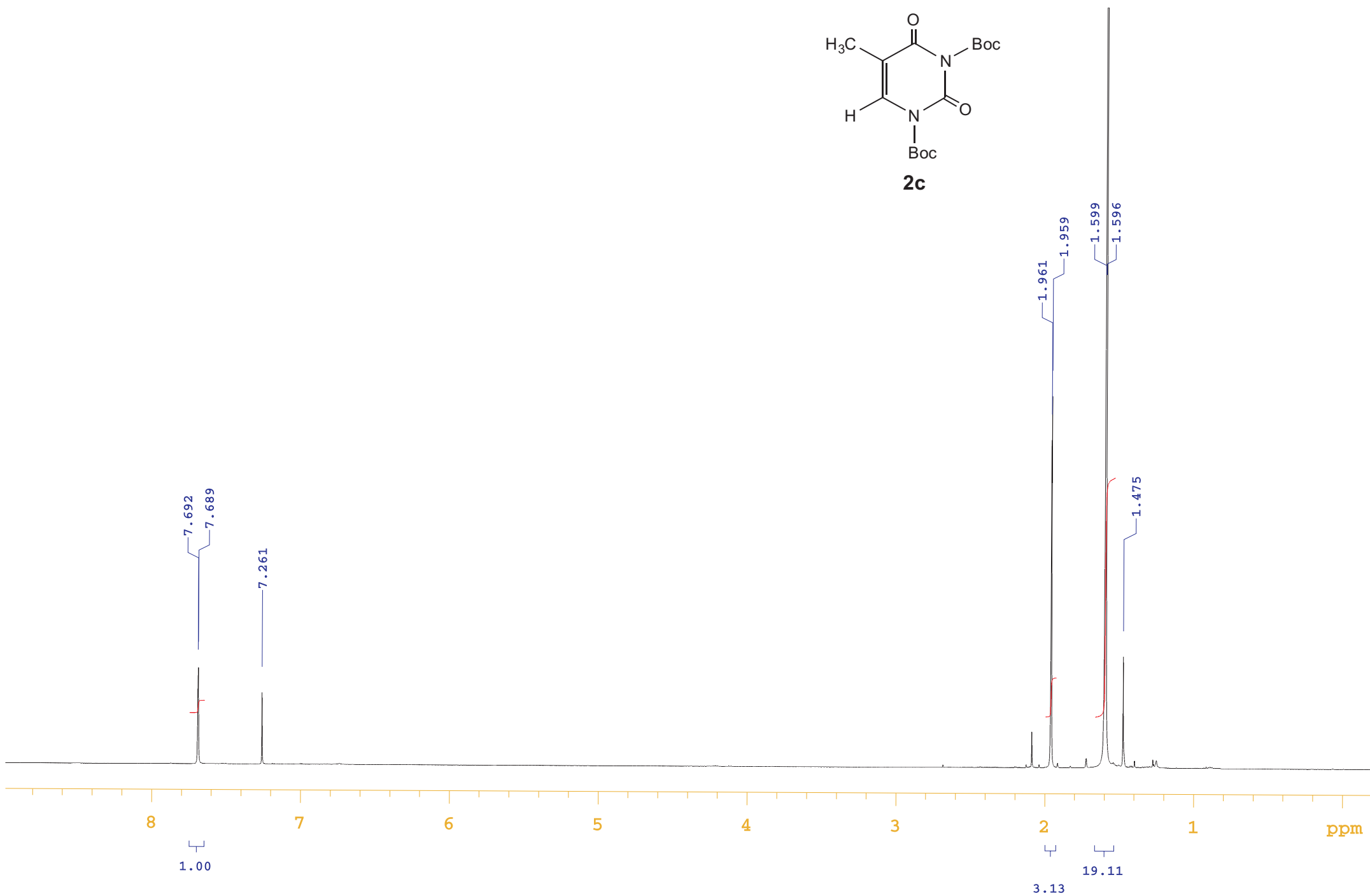
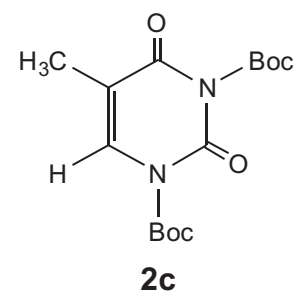
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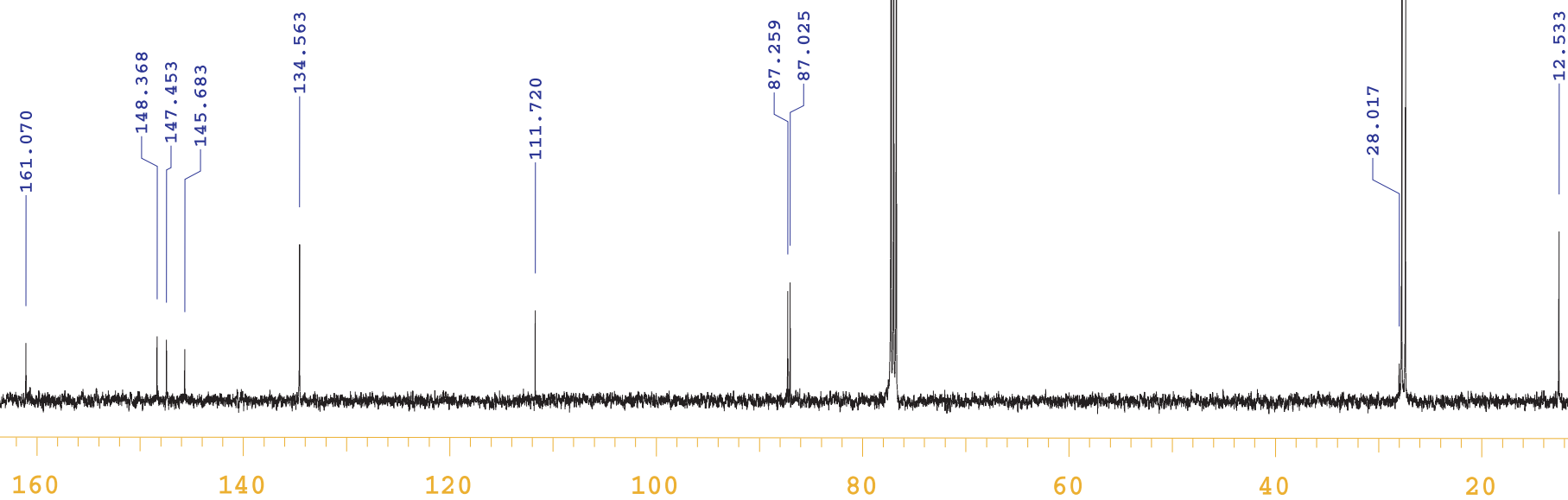
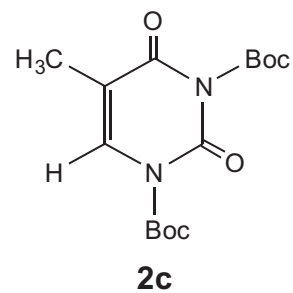


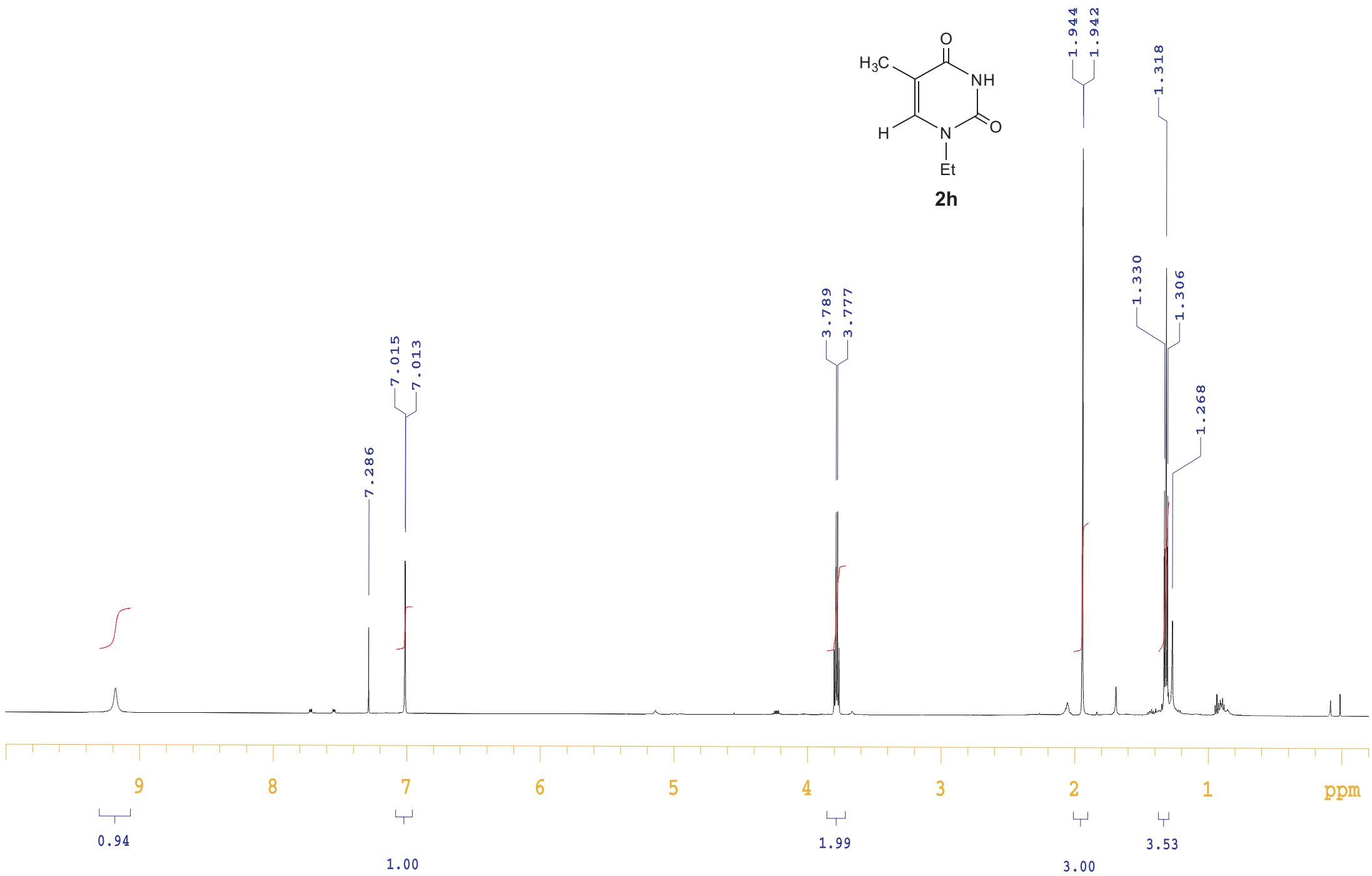
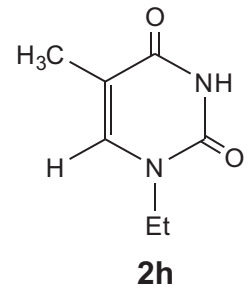


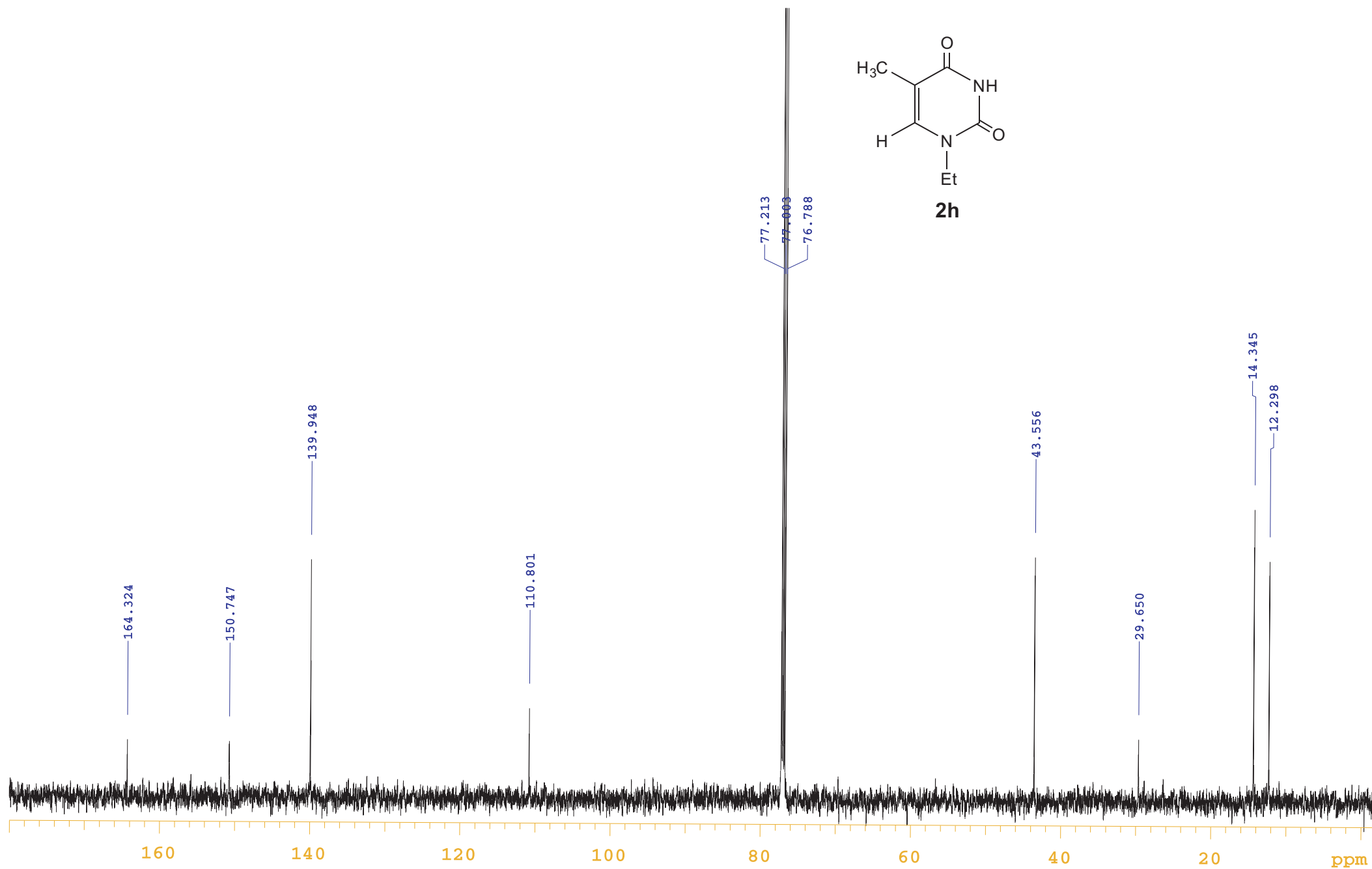
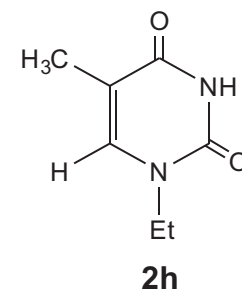
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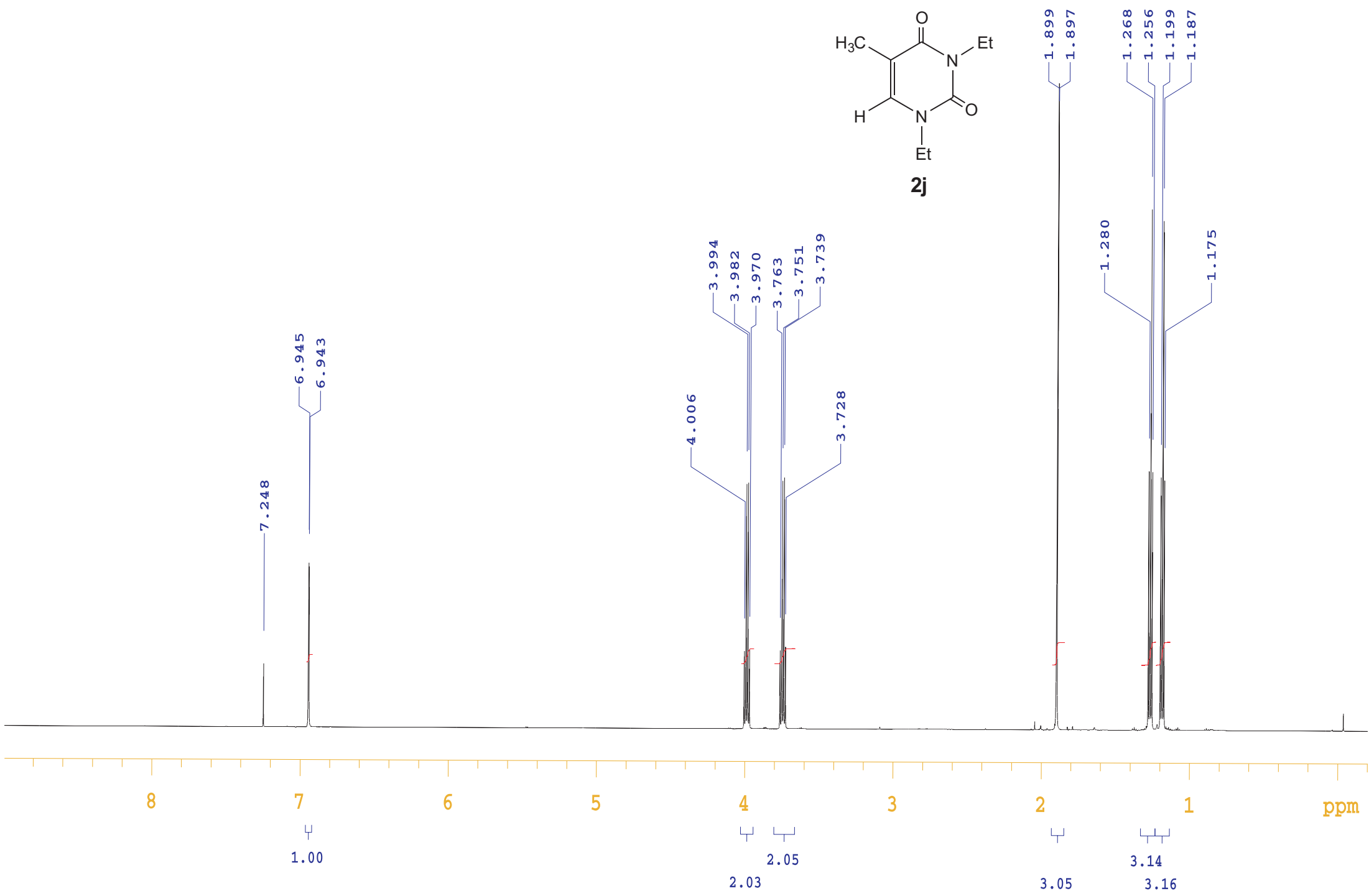
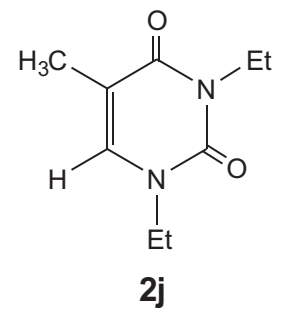


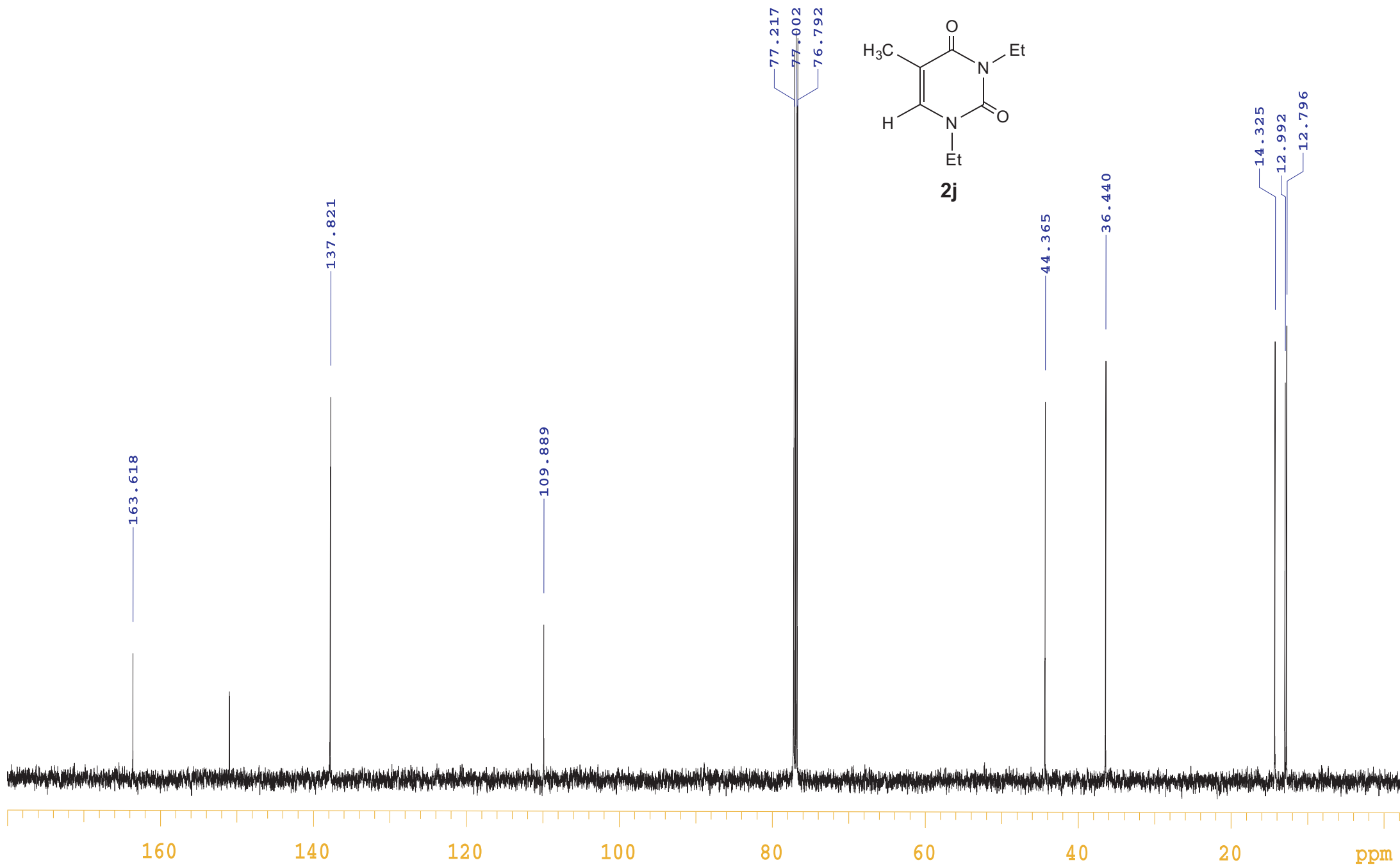


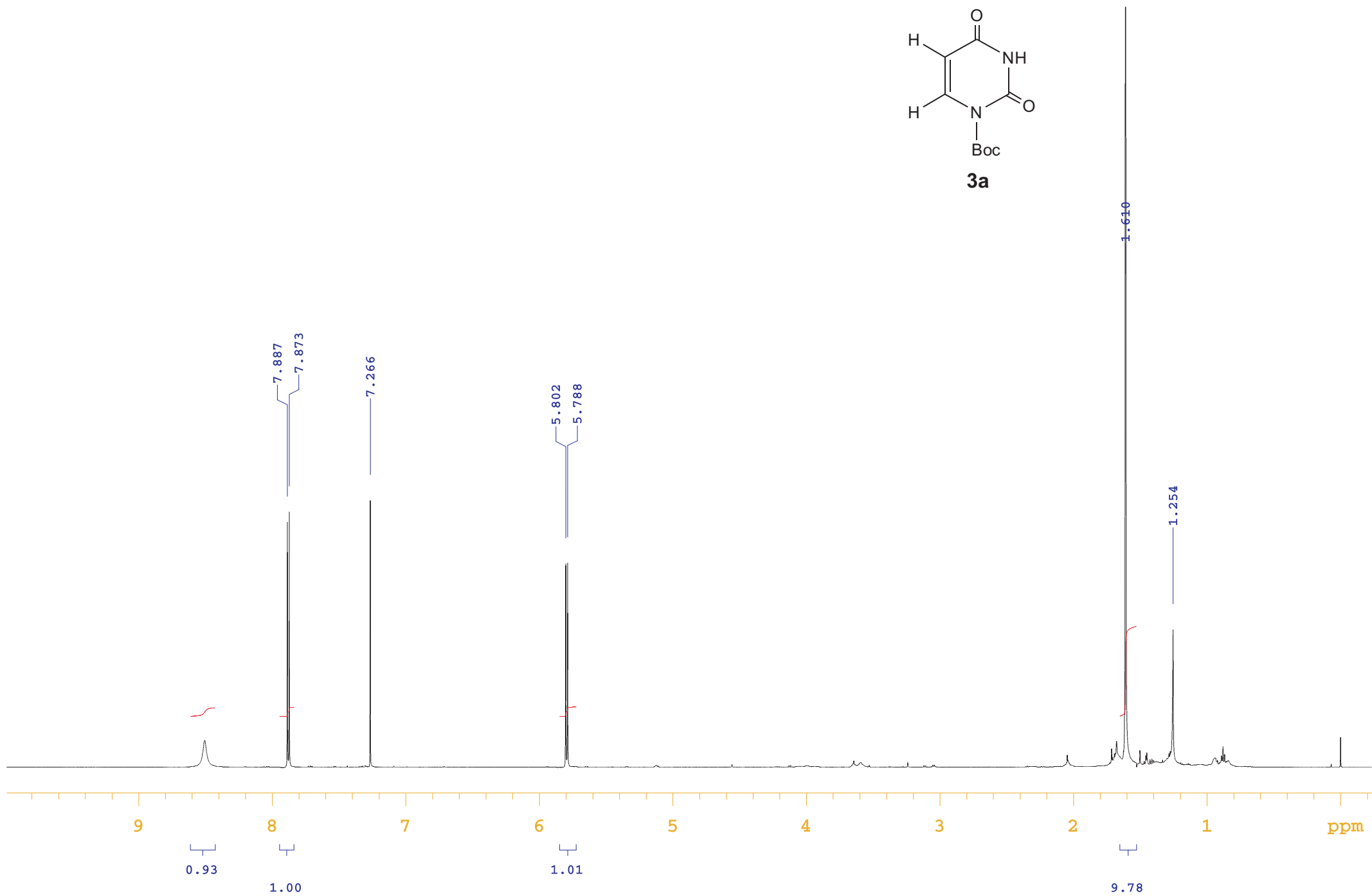
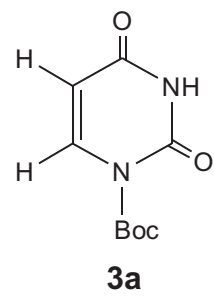


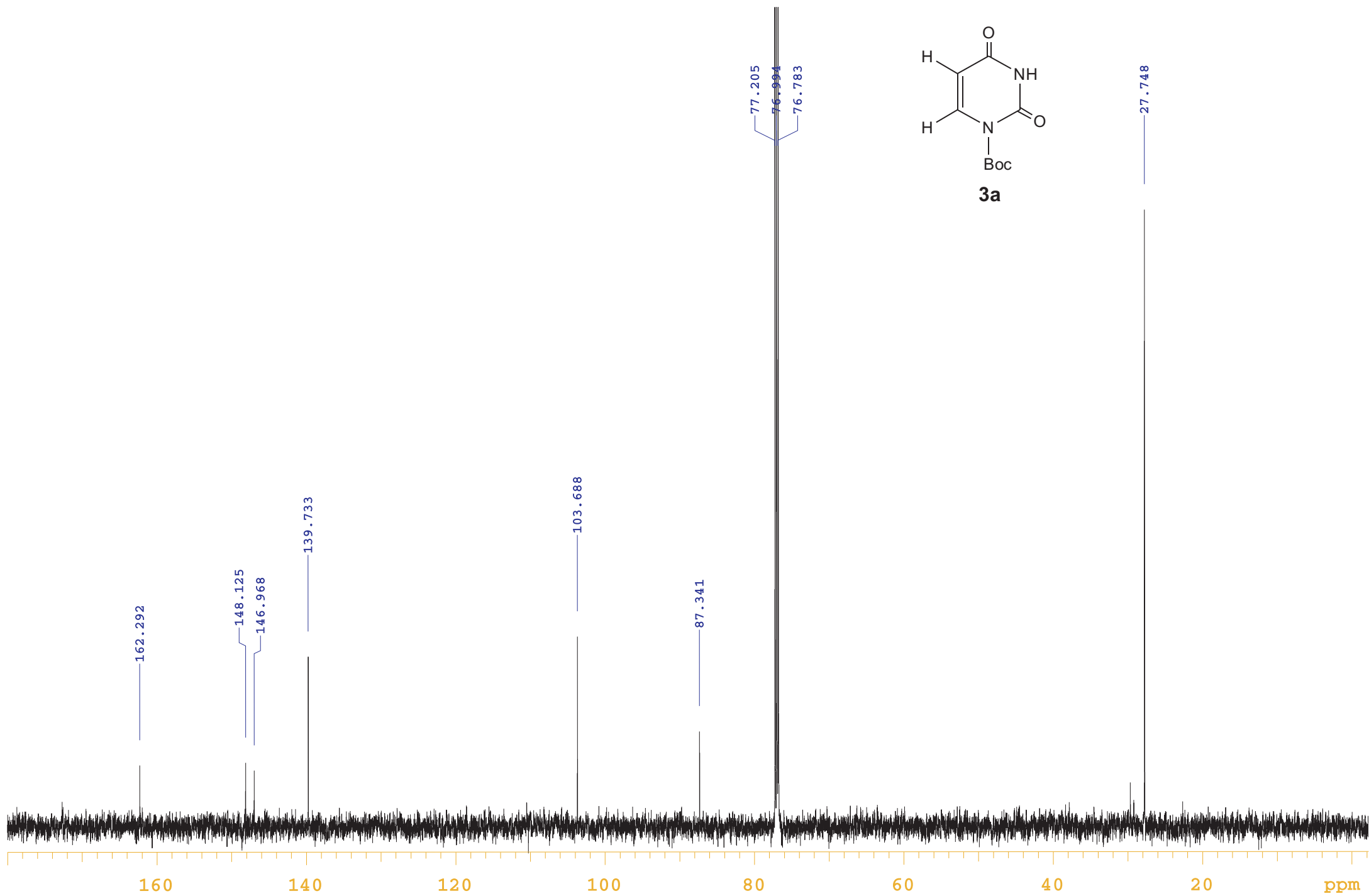


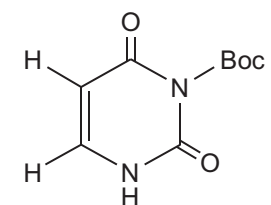




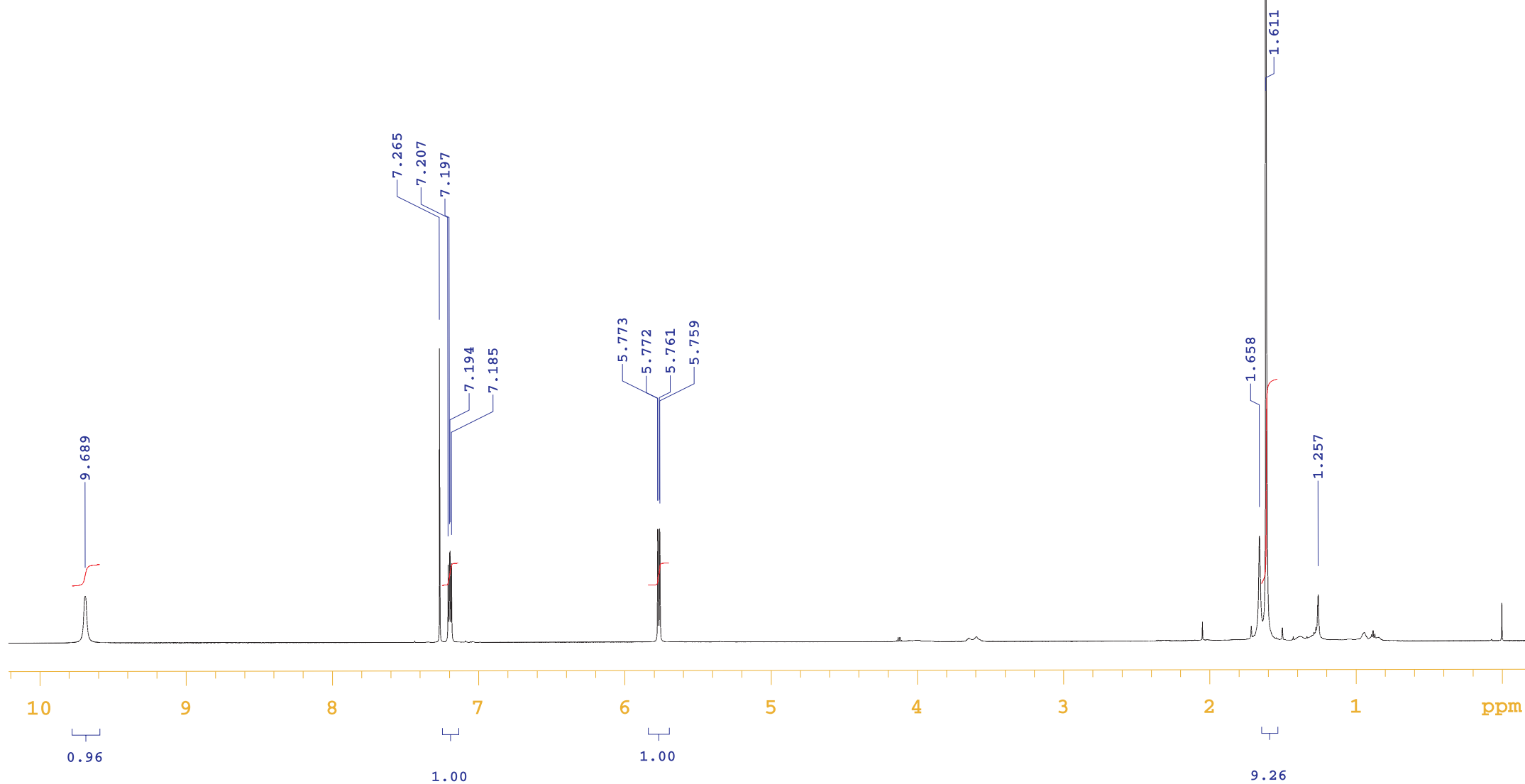


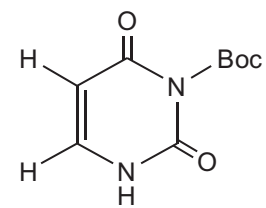




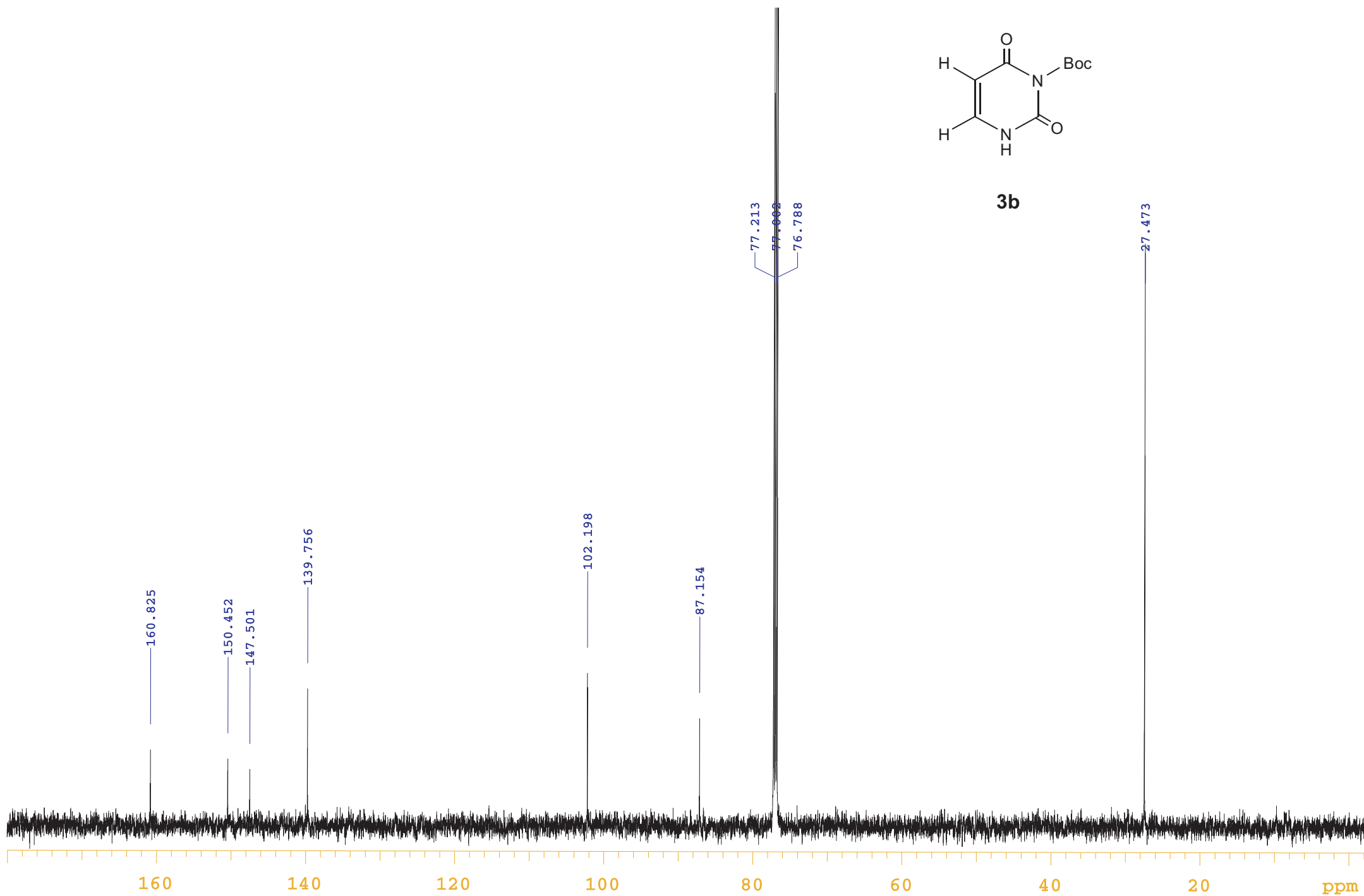


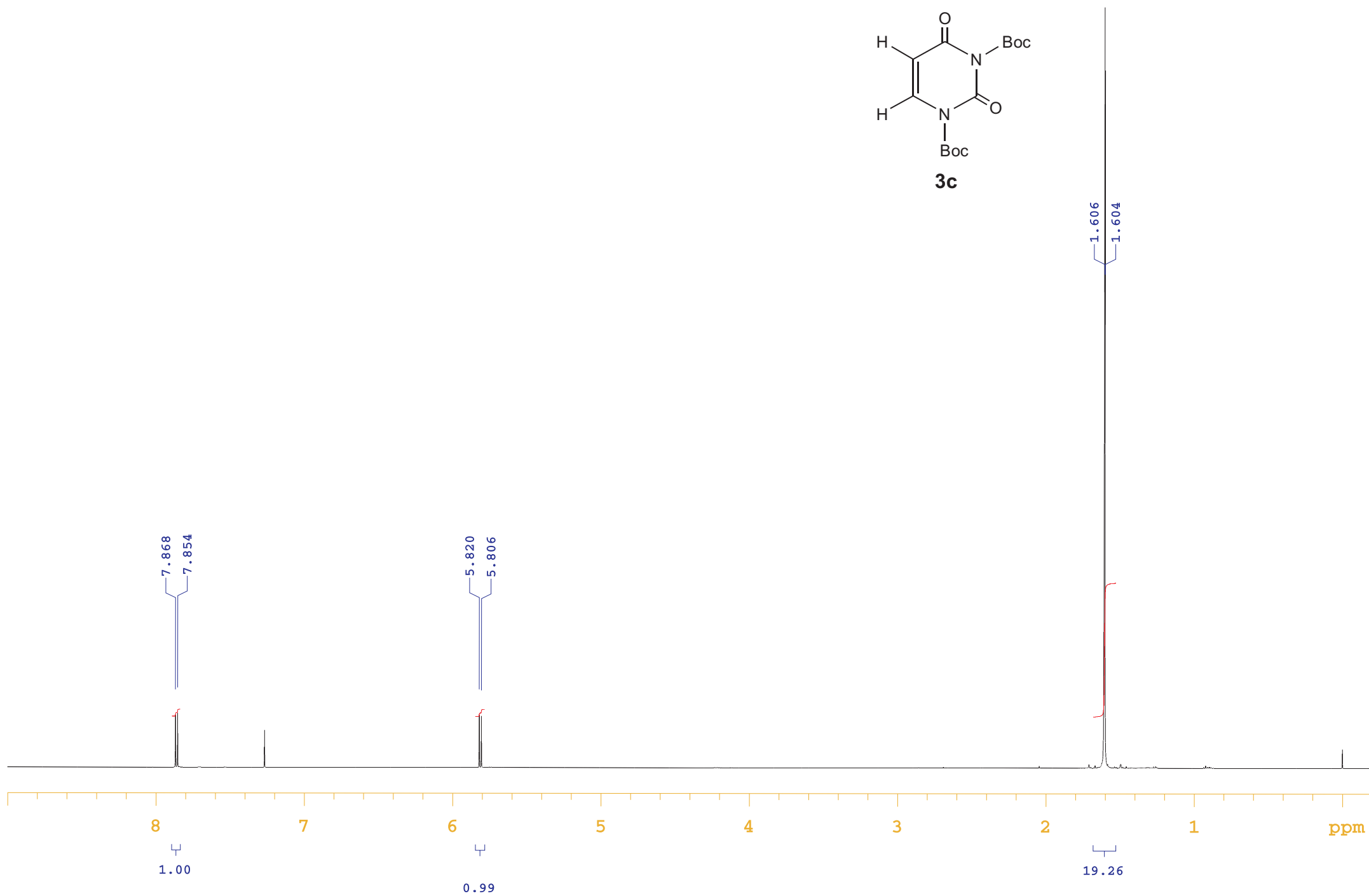
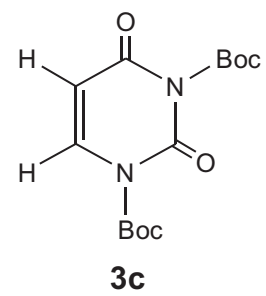
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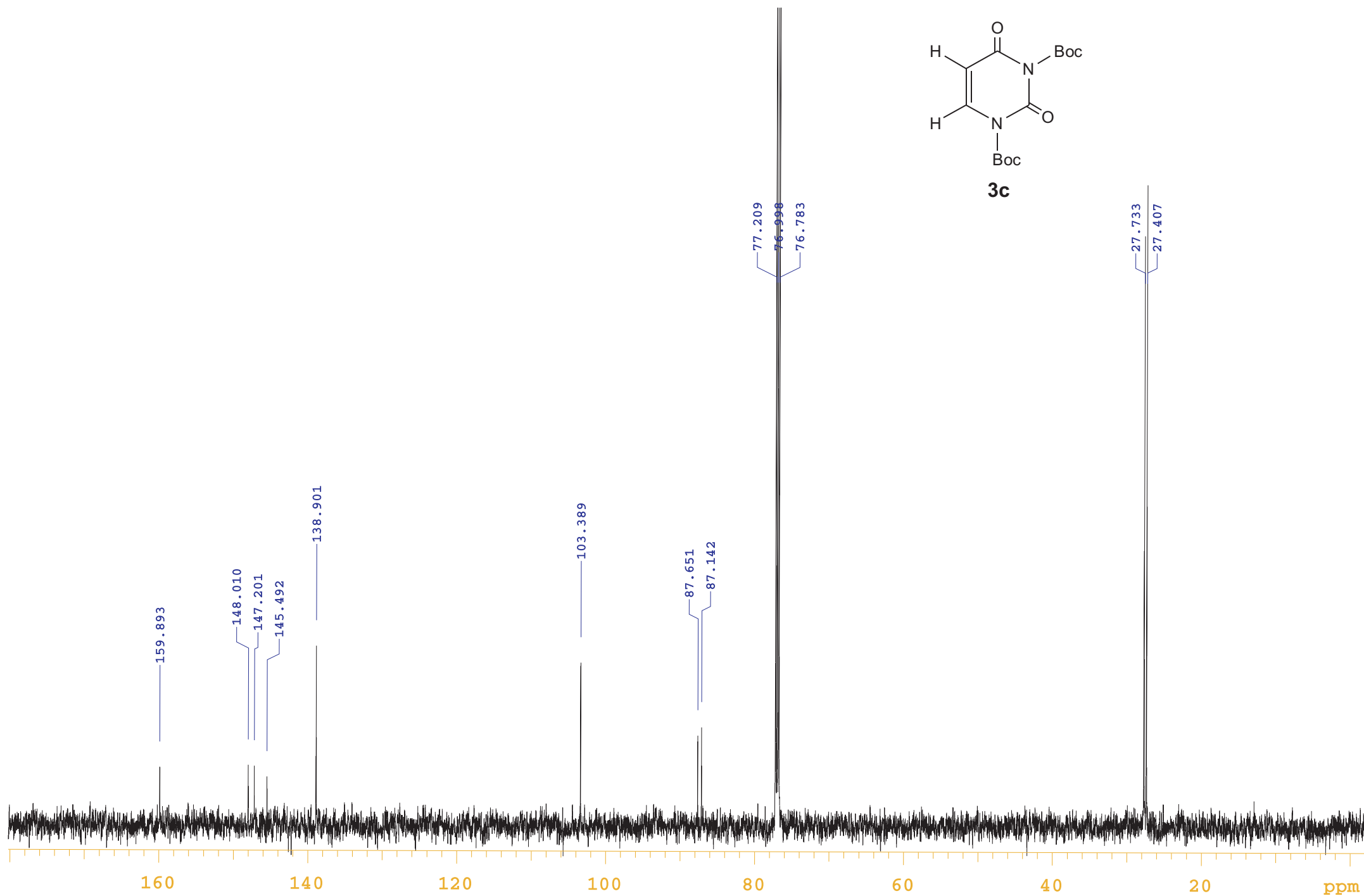
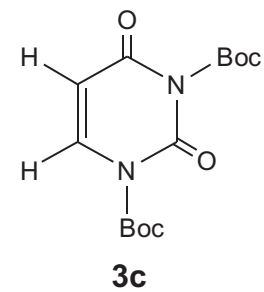


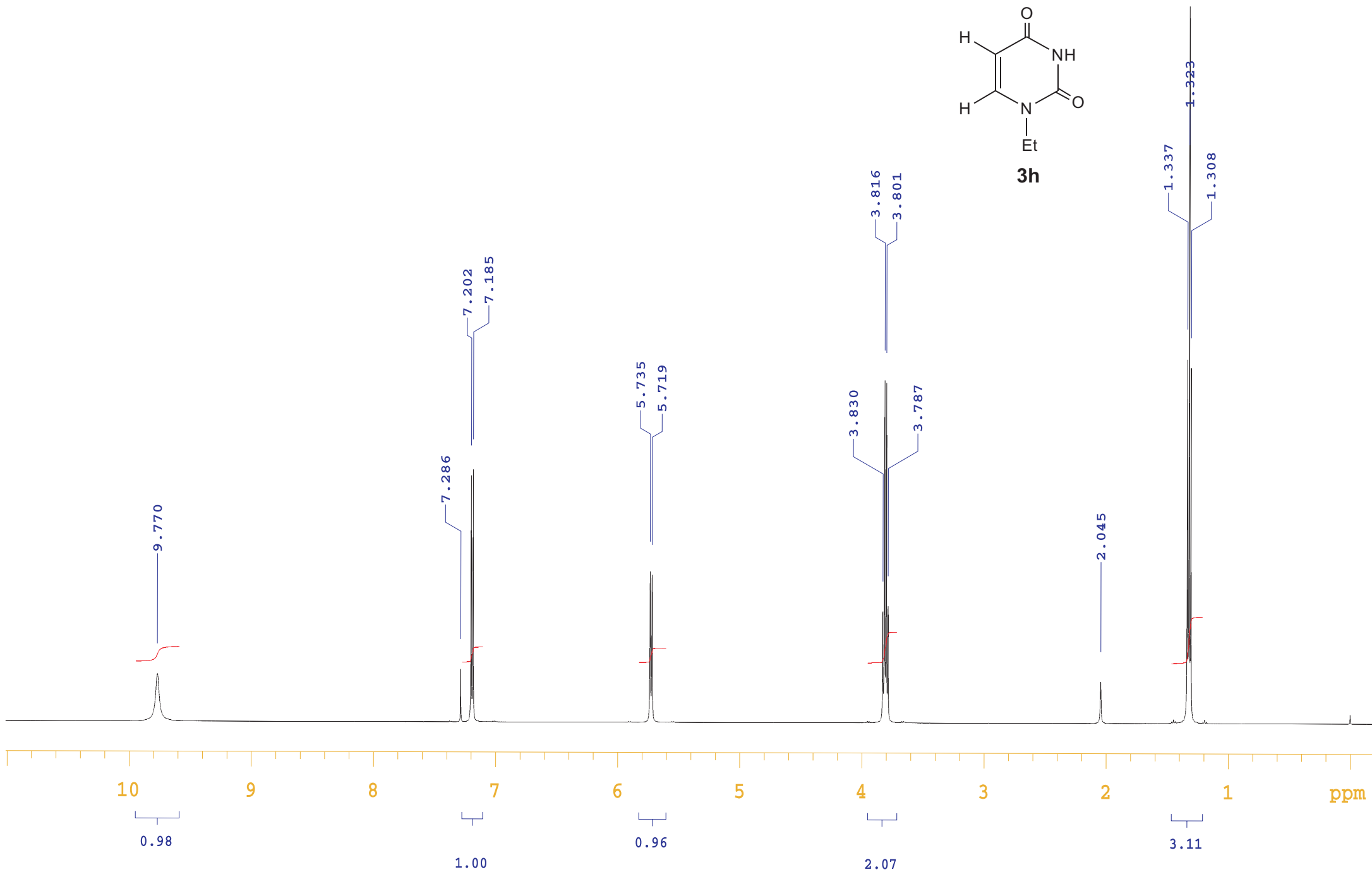
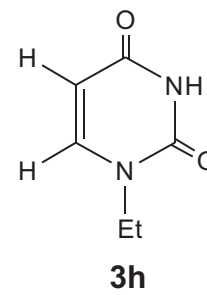


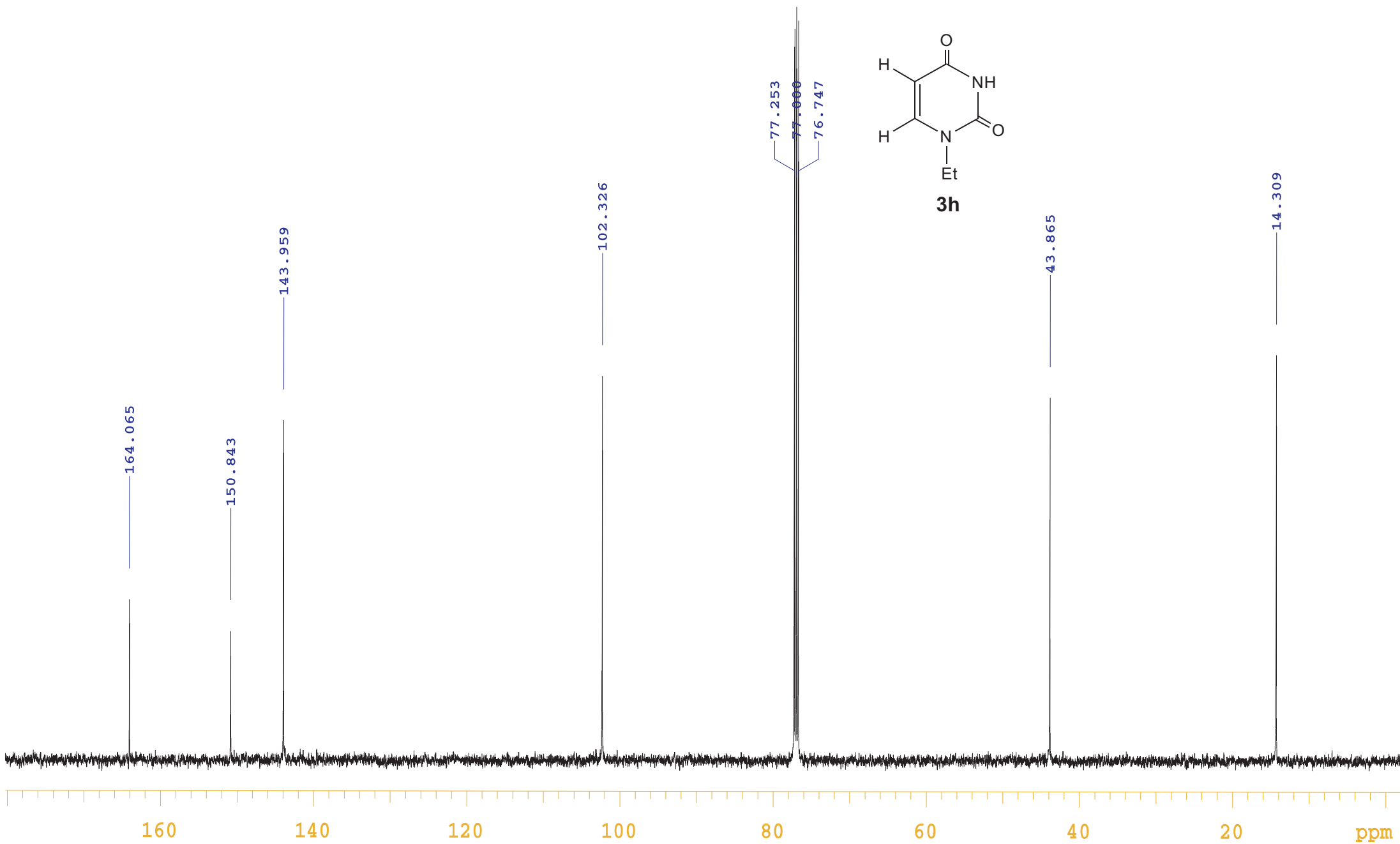
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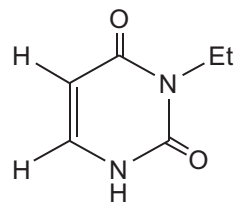




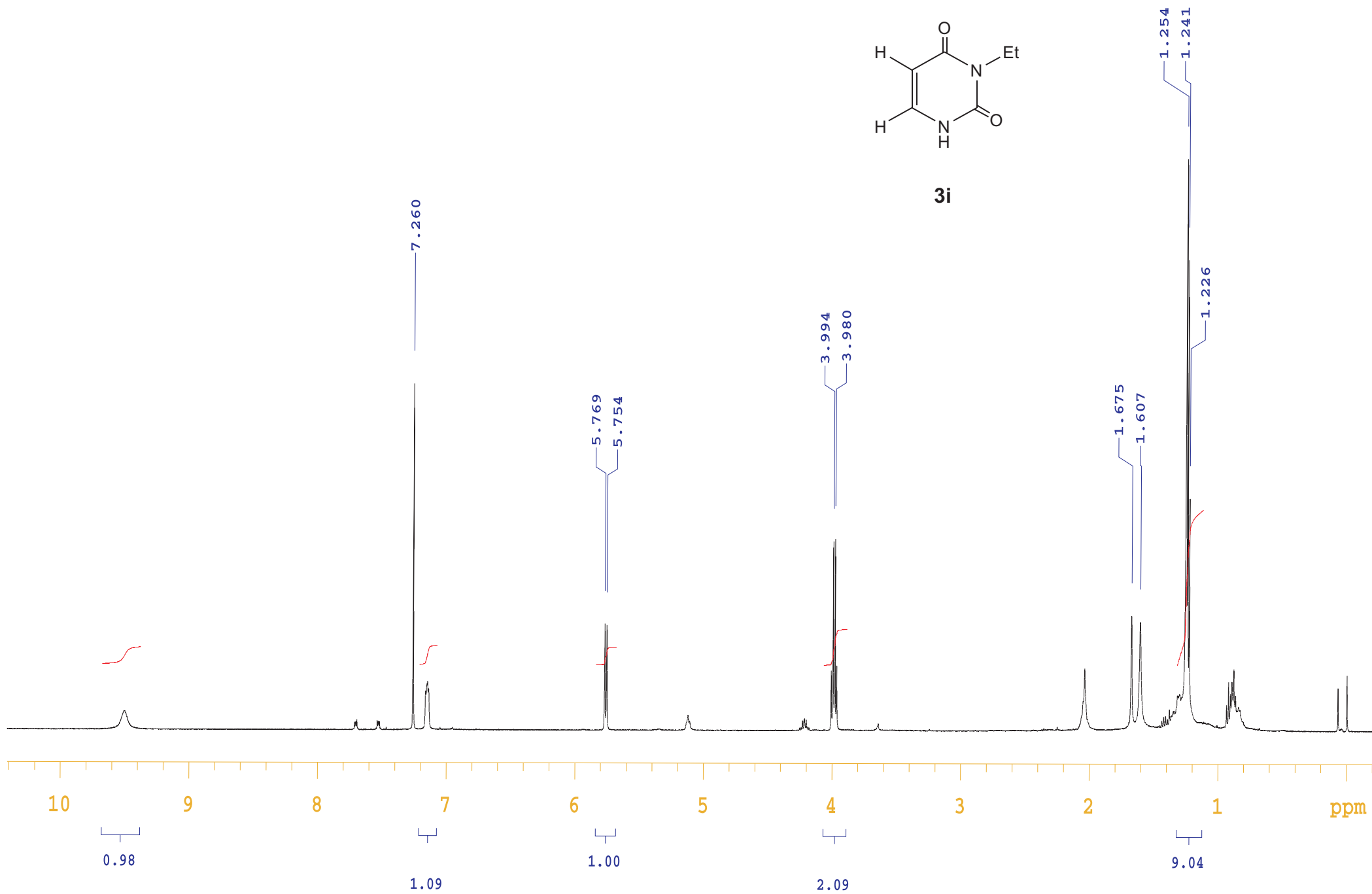


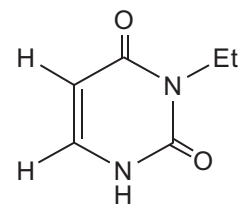






3i





3i

