

Metal-free TEMPO Promoted C(sp³)-H Amination to Afford Multisubstituted Benzimidazoles

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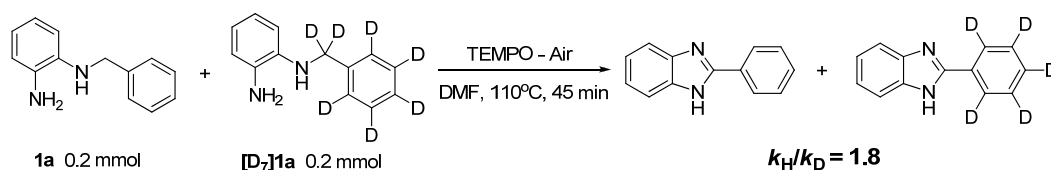
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Supporting Information

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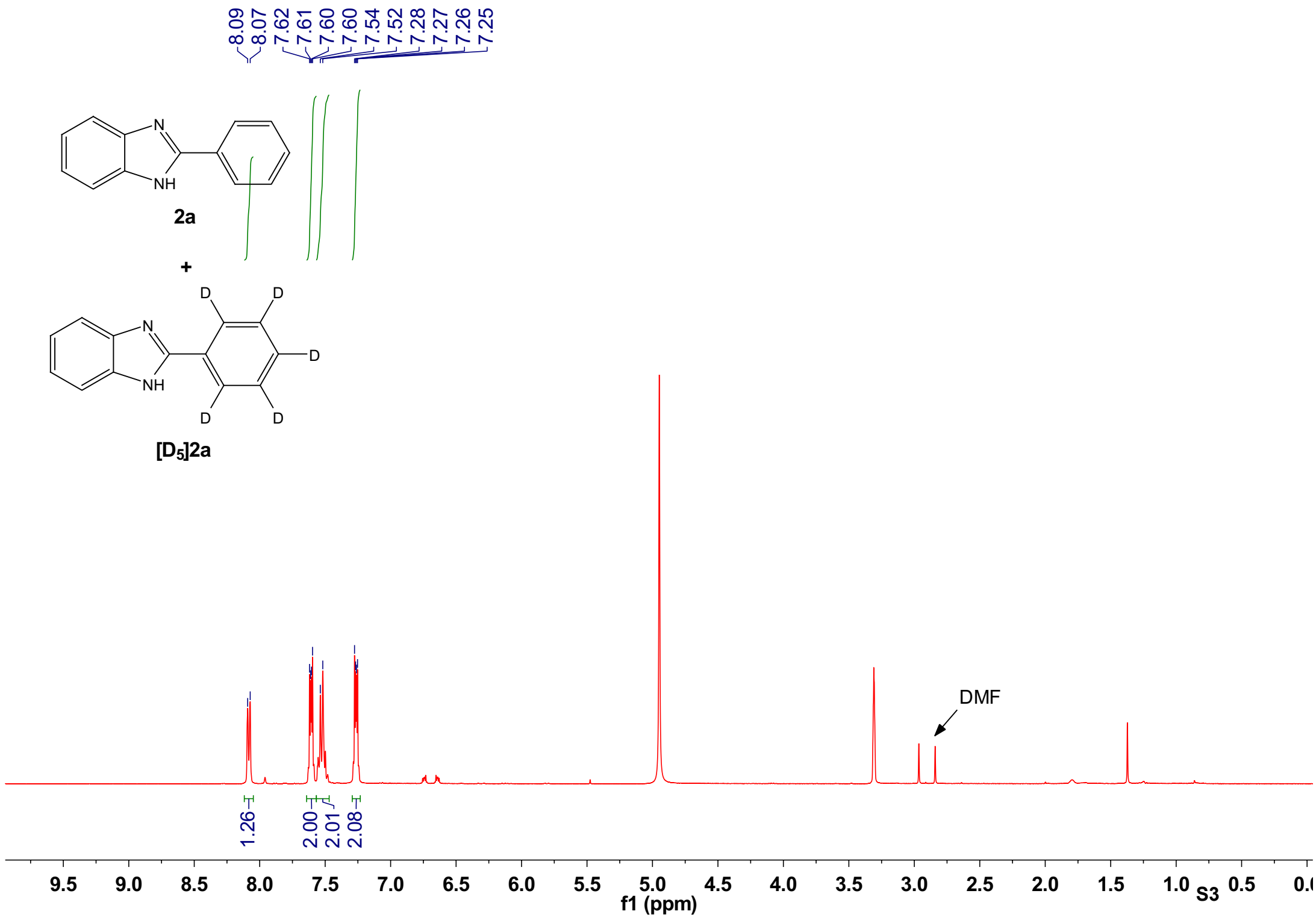
Studies on the kinetic isotope effects

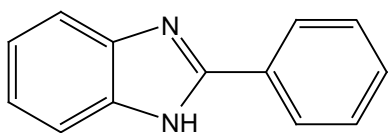


To a solution of **1a** (40 mg, 0.2 mmol) and [**D**₇]**1a** (41 mg, 0.2 mmol) in DMF (4 mL) was added TEMPO (125 mg, 0.8 mmol), and the mixture was stirred at 110 °C for 45 min. The mixture was then diluted with AcOEt, washed with water and saturated NaCl aq. solution, then dried over anhydrous Na₂SO₄. After filtration, the filtrate was concentrated in vacuo. The residue was purified by silica gel column chromatography and subsequently washed by n-hexane. The pure mixture of products was characterized by ¹H NMR. Kinetic isotope effect $k_H/k_D = 1.8$ was determined by taking the difference in the integration of protons from the 2-phenyl group of the benzimidazole product. The integration of the peak at 8.07-8.09 ppm was 1.26 instead of 1.99 and the peak at 7.48-7.55 ppm was 2.01 instead of 3.09.^{1,2}

References:

- (1) Matcha, K.; Antonchick, A. P. *Angew Chem Int Edit* **2013**, 52, 2082.
- (2) Yan, Y. Z.; Zhang, Y. H.; Feng, C. T.; Zha, Z. G.; Wang, Z. Y. *Angew Chem Int Edit* **2012**, 51, 8077.



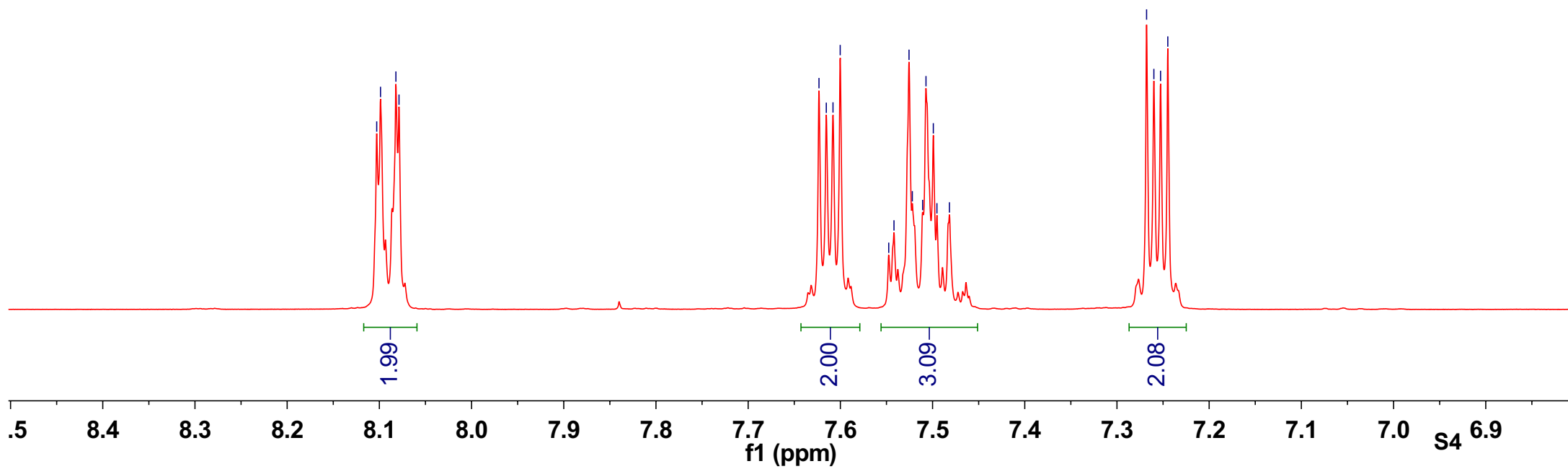


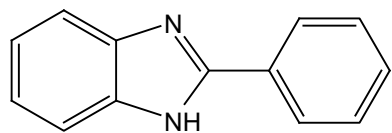
2a

8.10
8.10
8.08
8.08

7.62
7.62
7.61
7.60
7.55
7.54
7.53
7.52
7.51
7.51
7.50
7.50
7.48

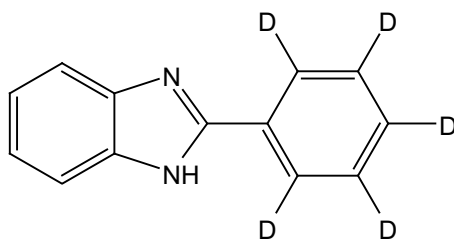
7.27
7.26
7.25
7.24





2a

+

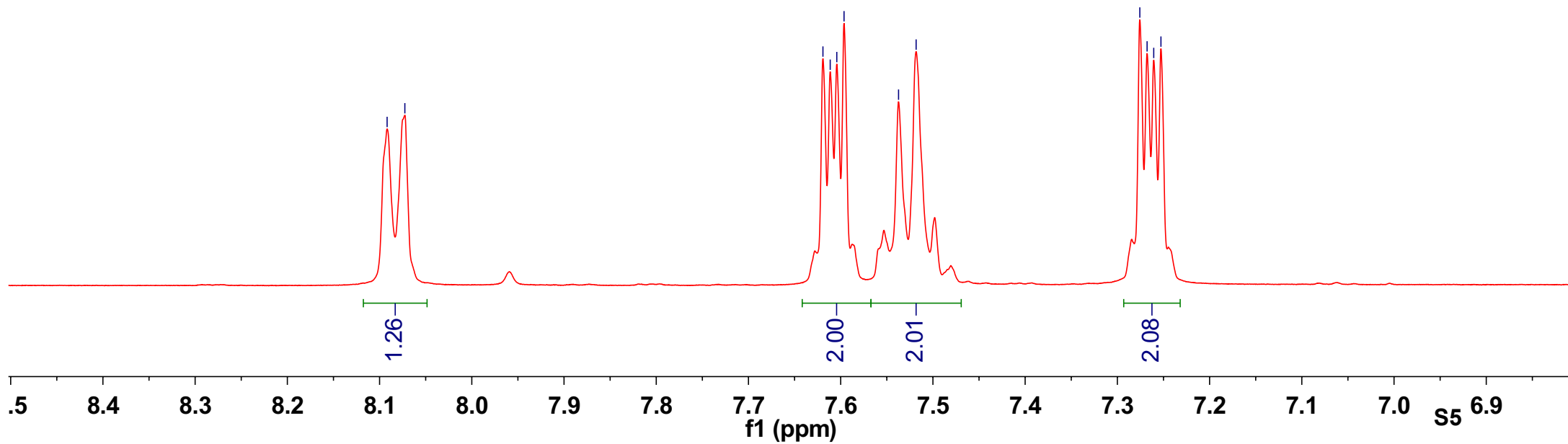


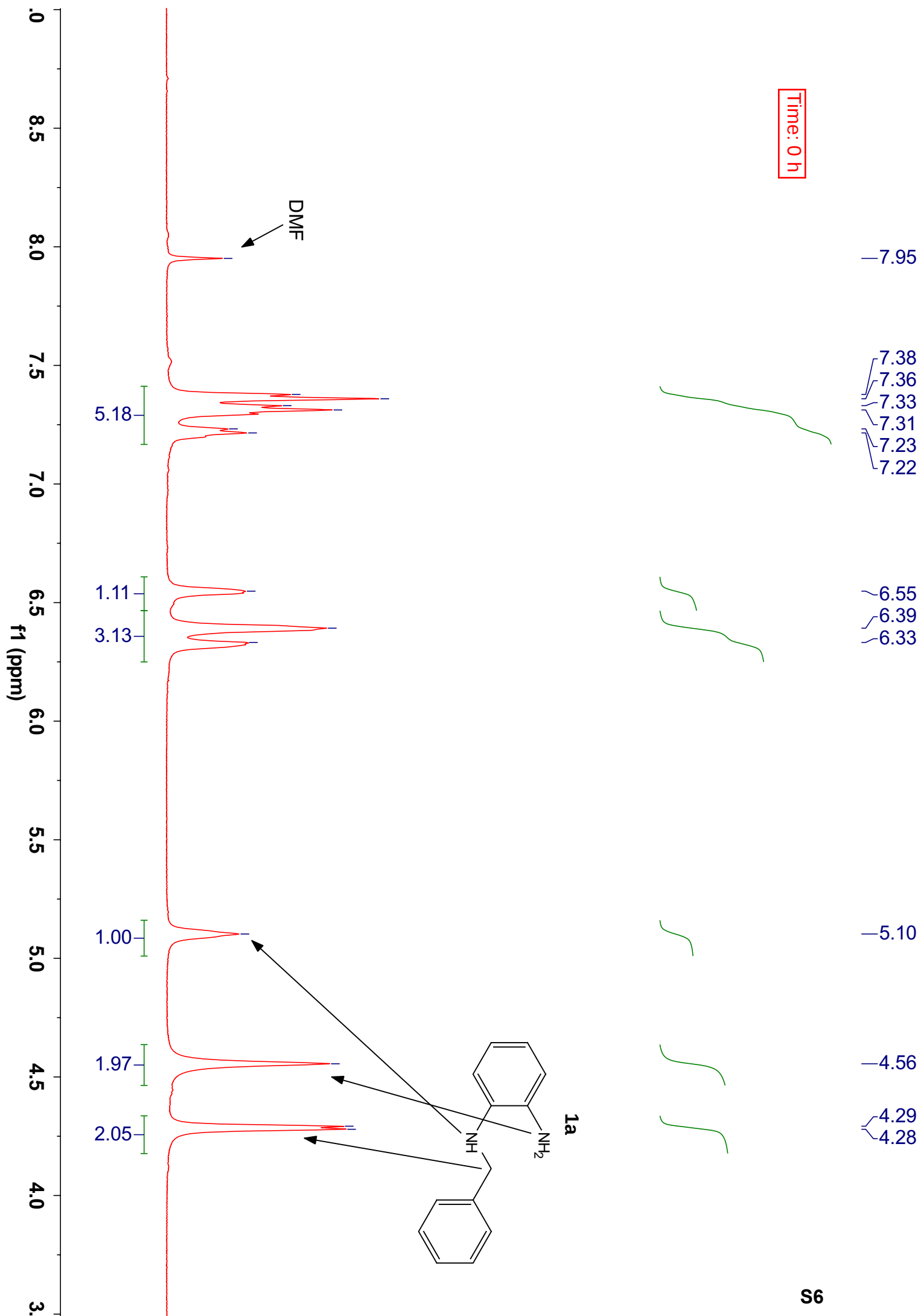
[D₅]2a

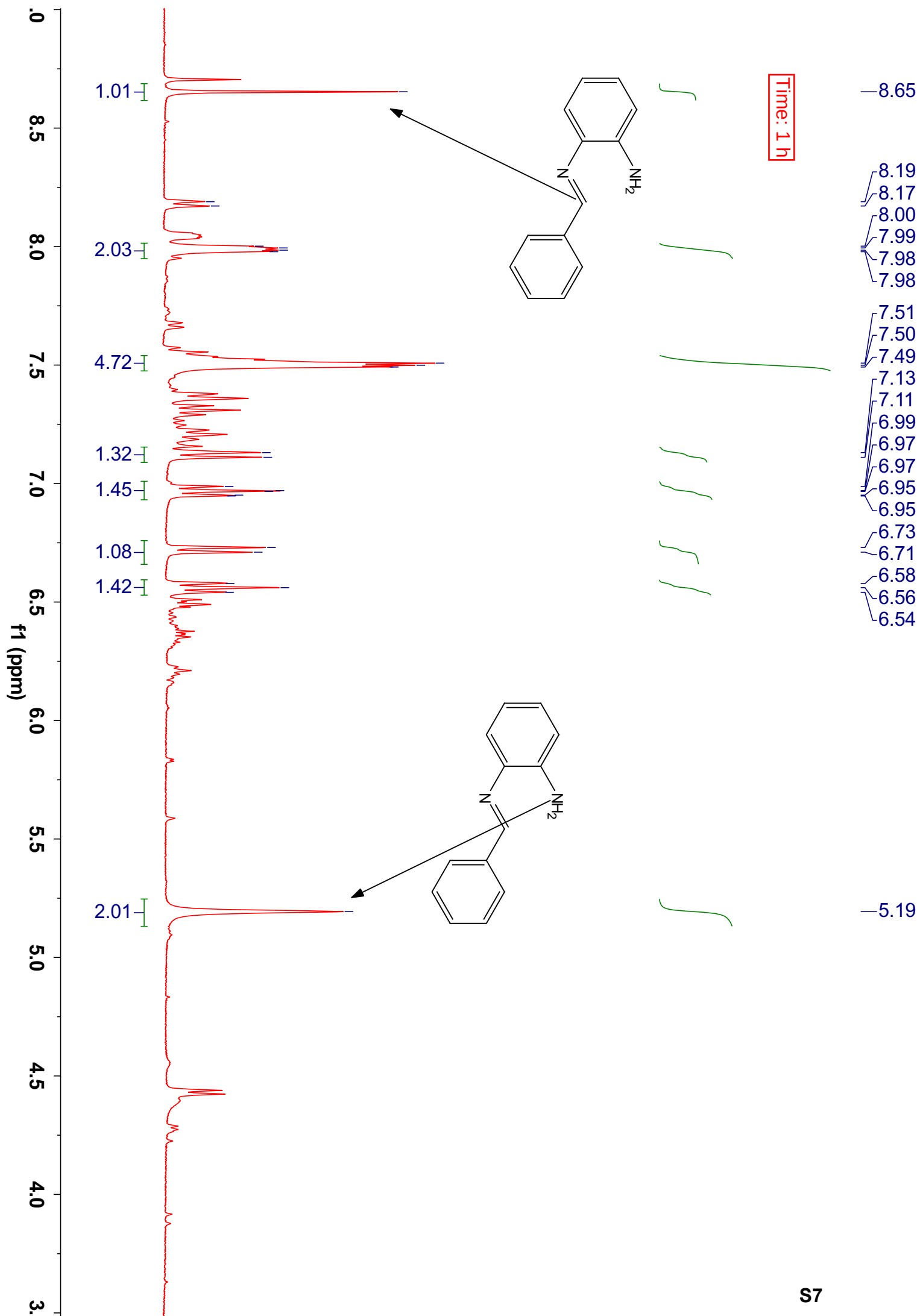
~8.09
~8.07

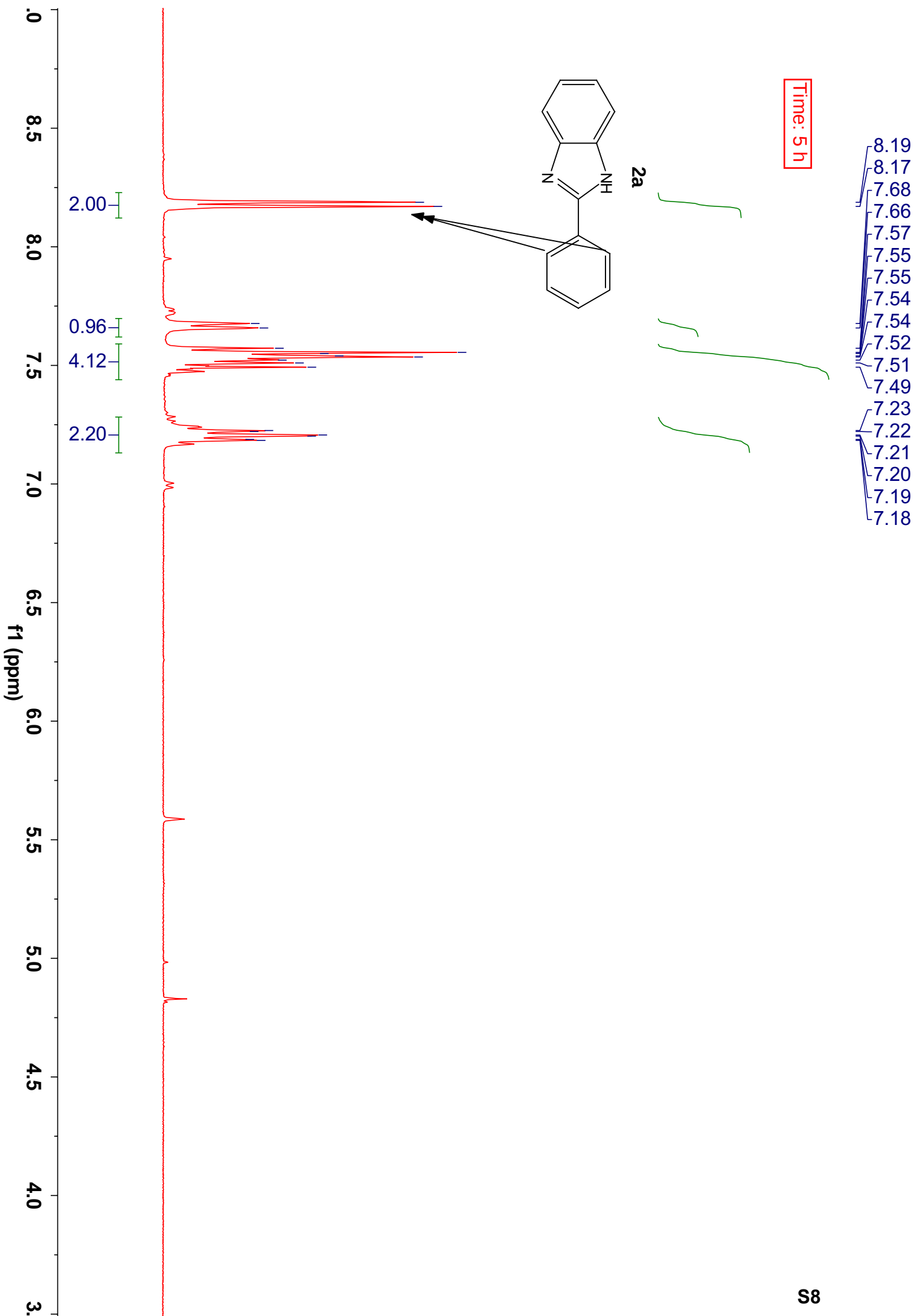
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7.28
7.27
7.26
7.25



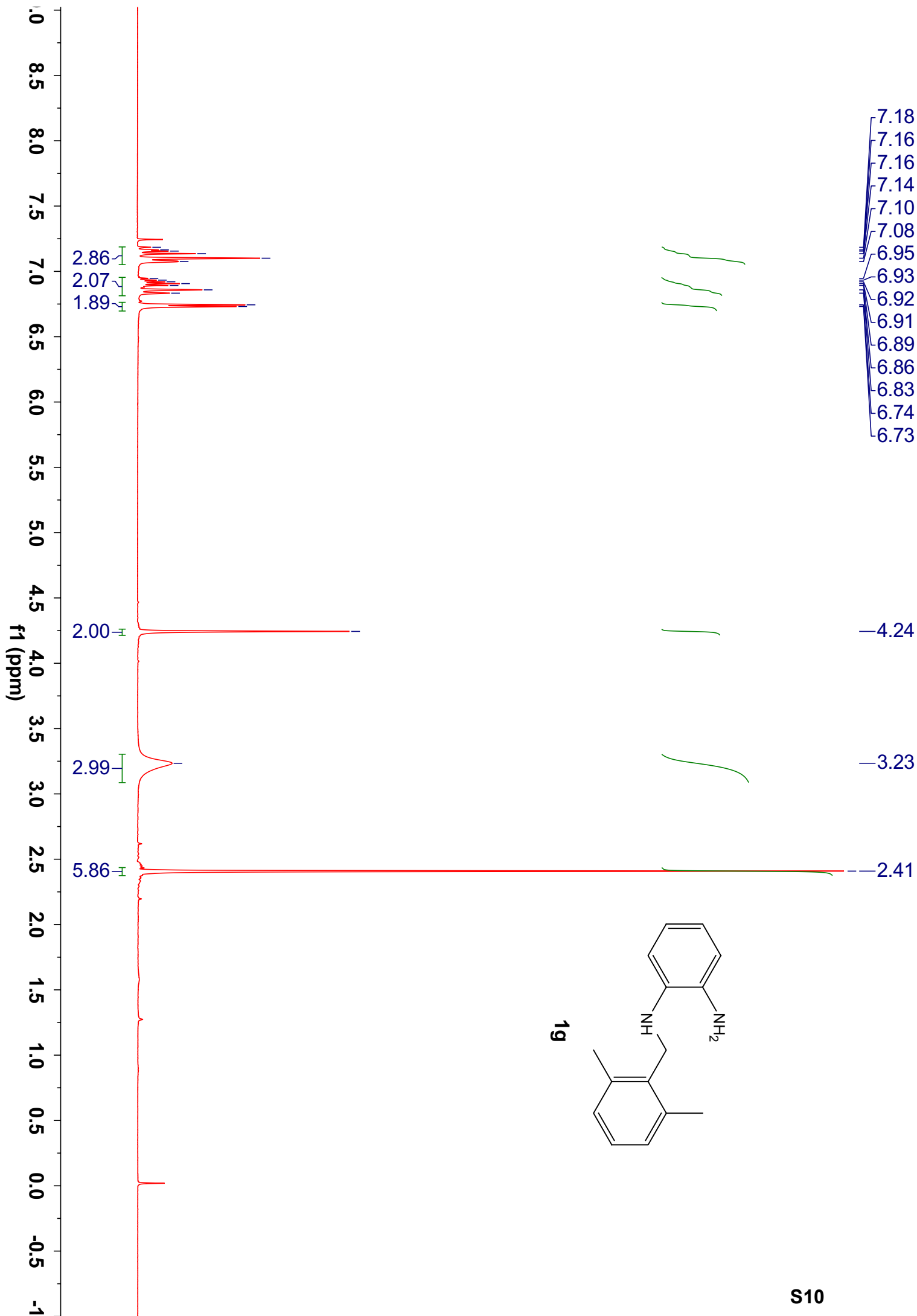


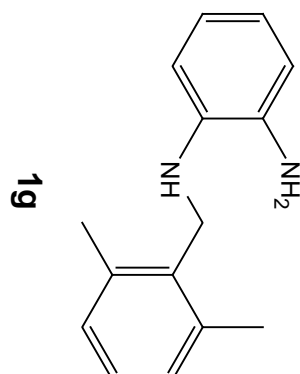




Notes:

- a. The reaction was carried out using method B (cat. TEMPO-O₂ condition) for the sufficient transformation of the starting material (**1a**) to the imine intermediate before the generation of benzimidazole product (**2a**).
- b. ¹H NMR Spectra were collected before the reaction started (0 h), when the starting material was completely converted to the intermediate determined by TLC (1 h) and when the reaction completed (5h).
- c. 300 μ l of the reaction solution was directly evaporated to dryness and rapidly redissolved by DMSO-d₆ for the ¹H NMR detection at specific time.

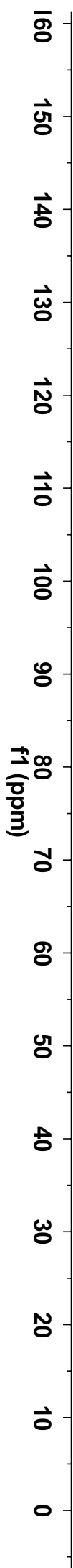


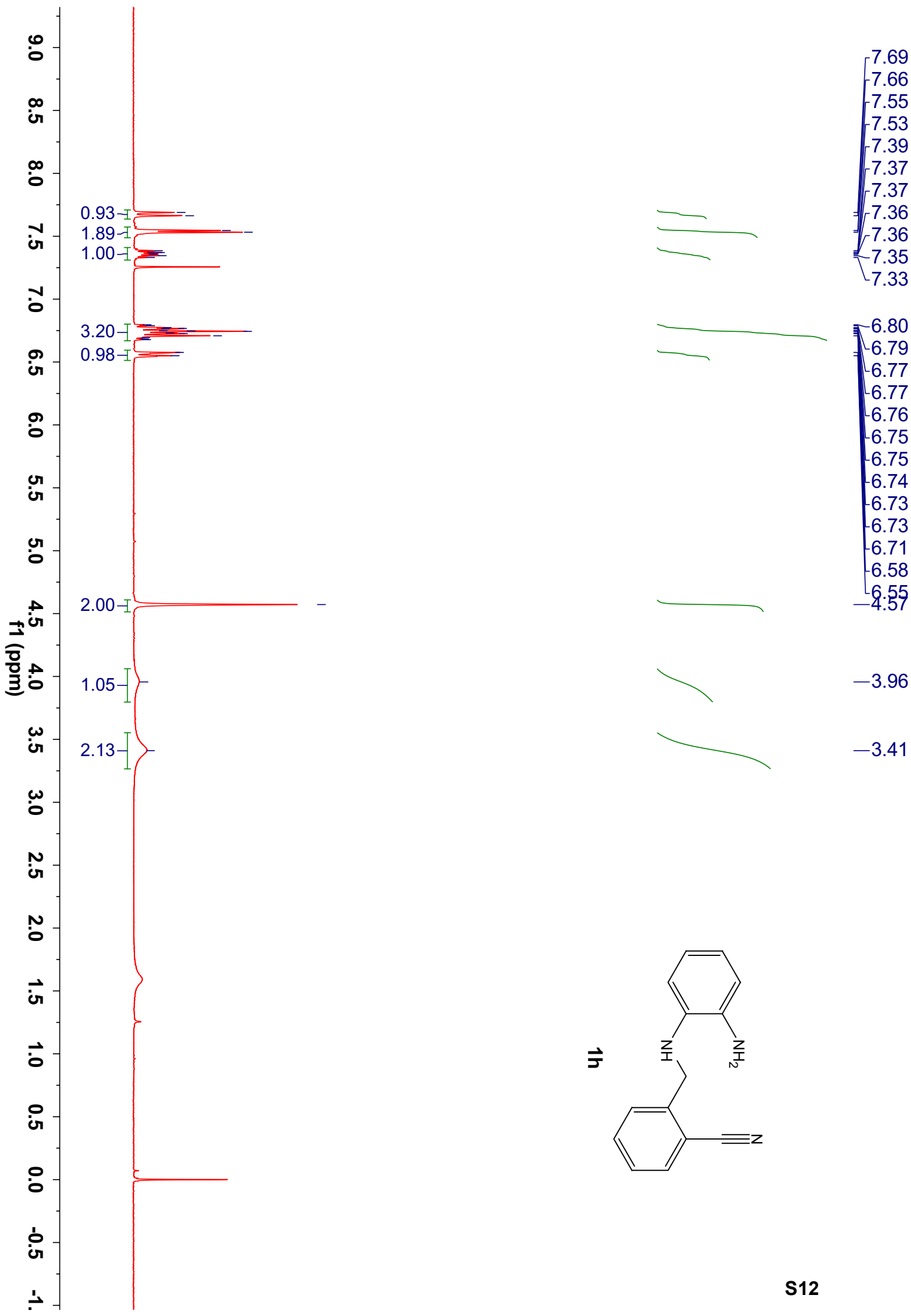
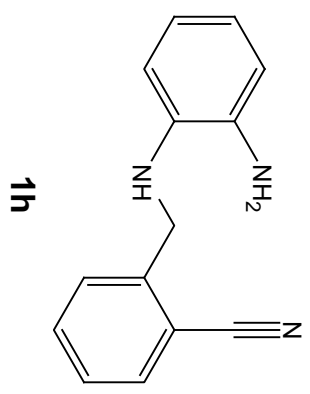


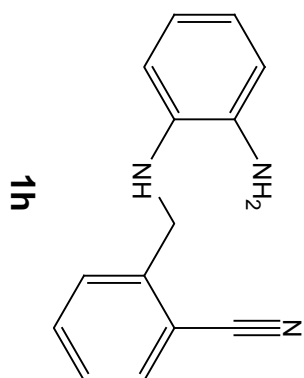
—42.92

—19.61

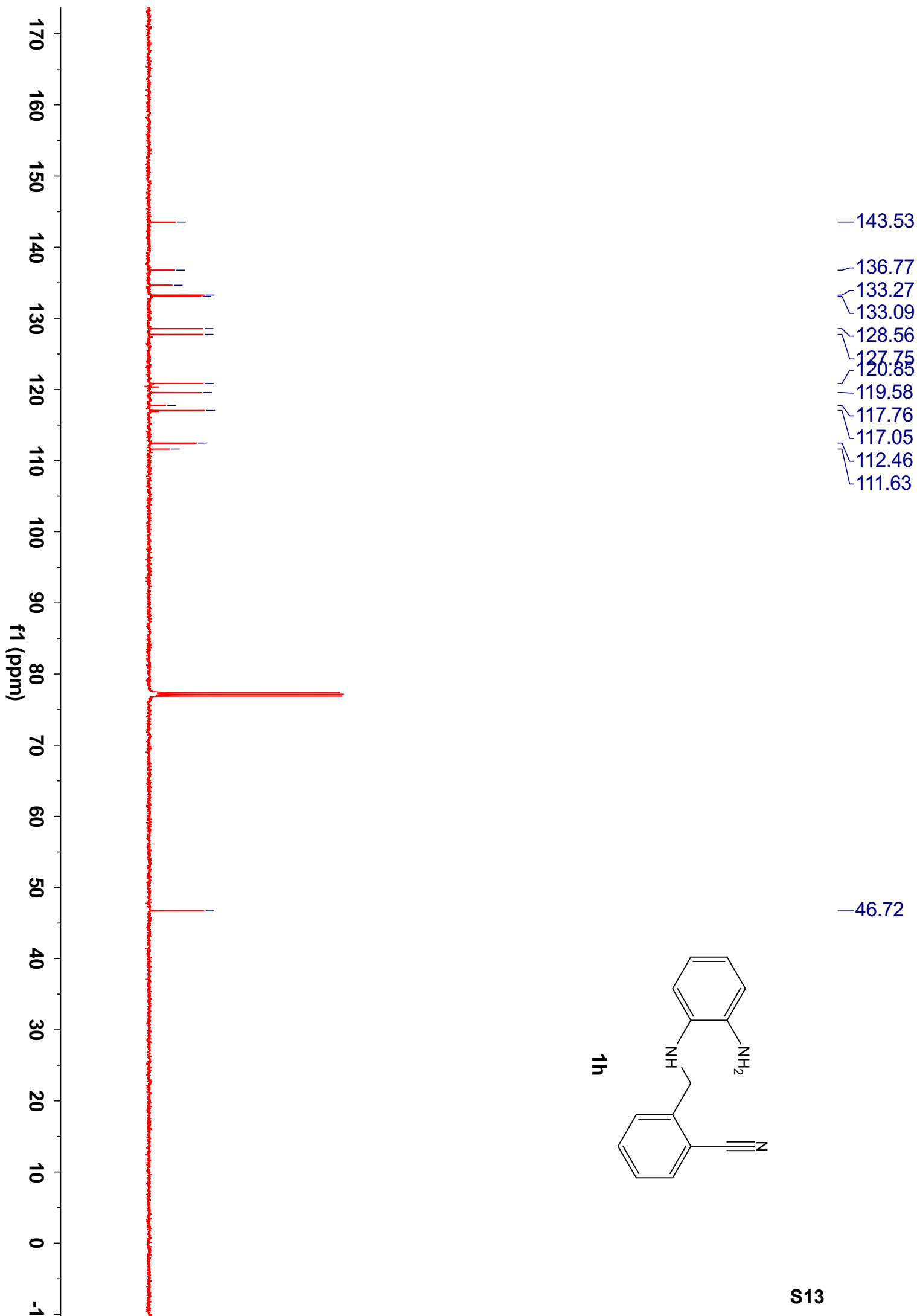
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137.81
135.48
134.34
128.44
127.74
120.78
118.73
116.34
111.48

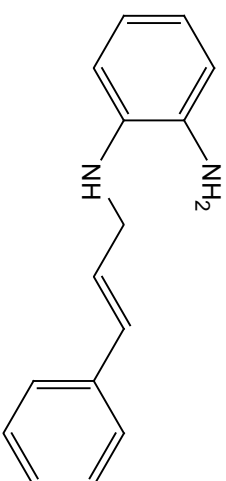






S13

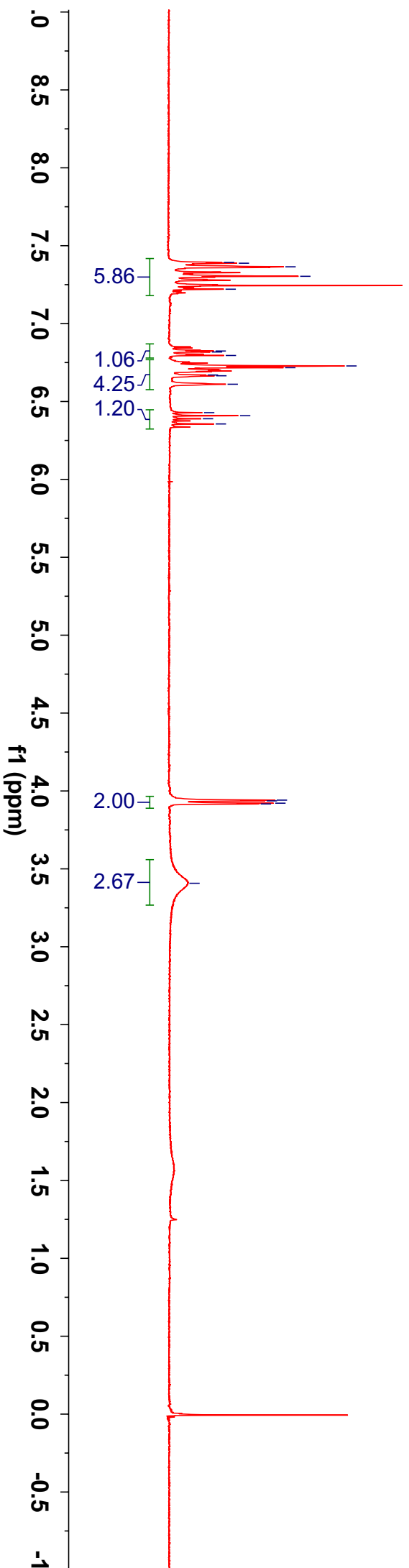


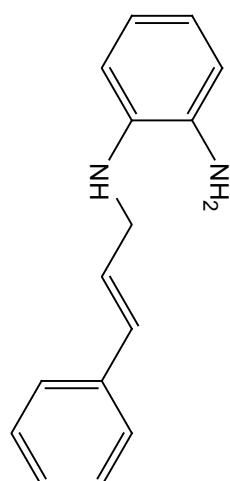


1j

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7.39
7.36
7.30
7.22
6.82
6.82
6.80
6.73
6.72
6.67
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6.41
6.39
6.36

3.94
3.94
3.92
3.92
3.41



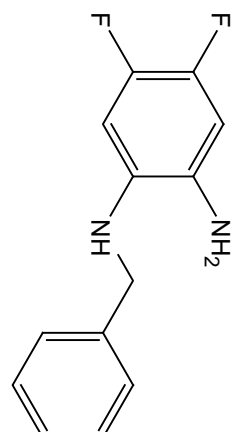
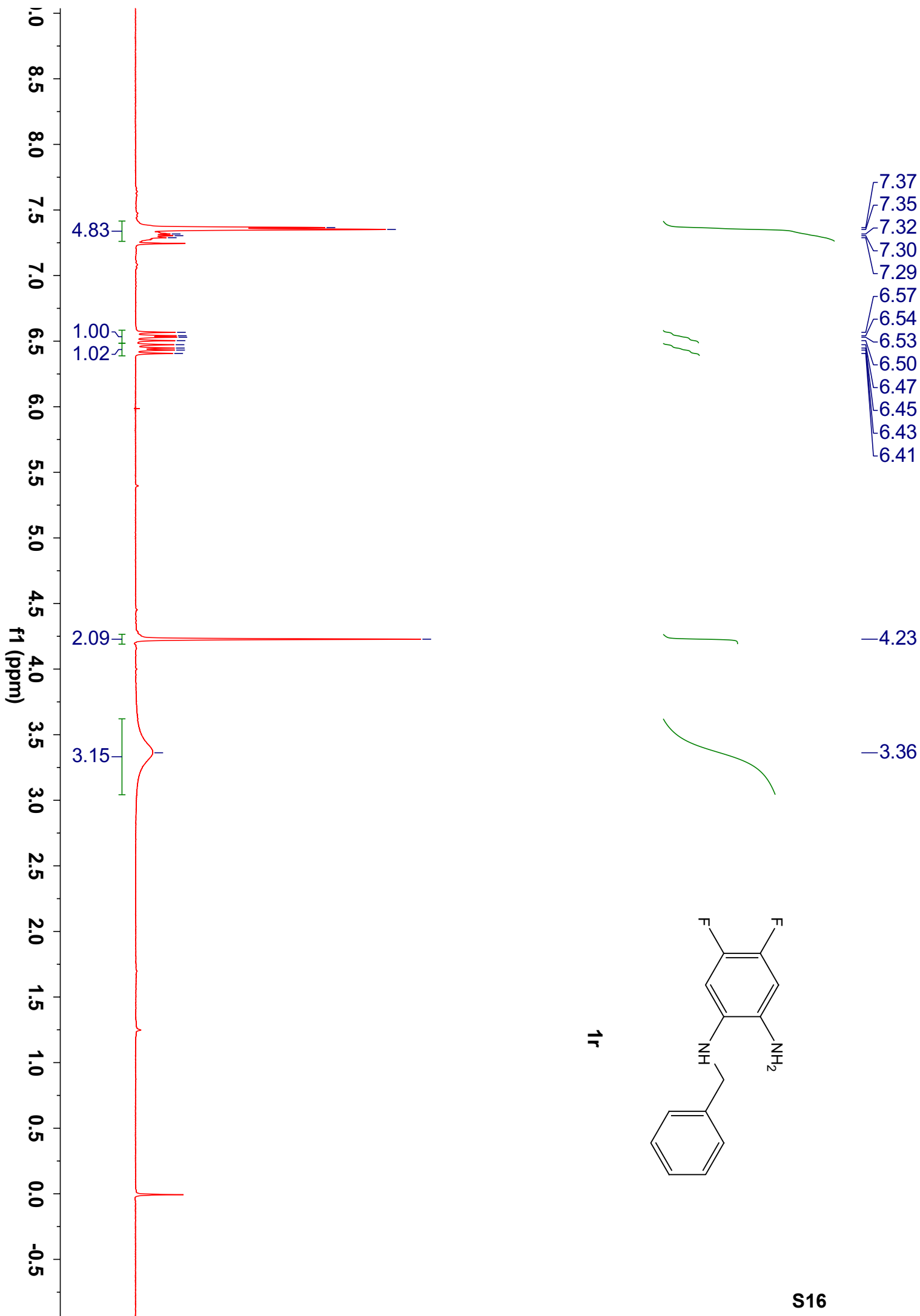


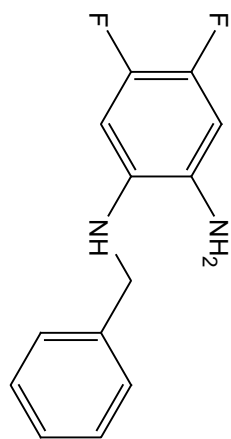
1j

—46.59

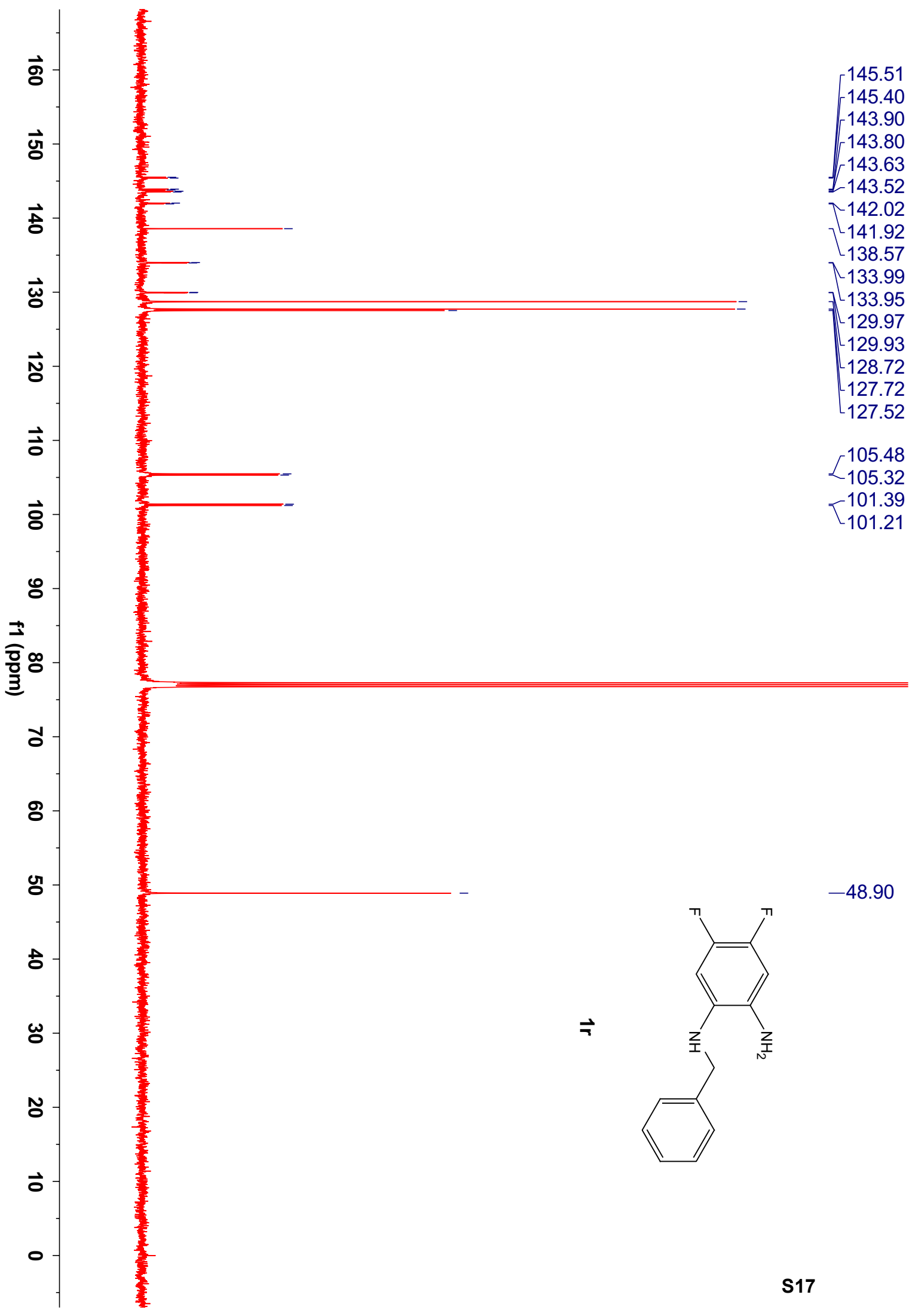
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131.74
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127.64
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126.45
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119.00
116.68
112.29

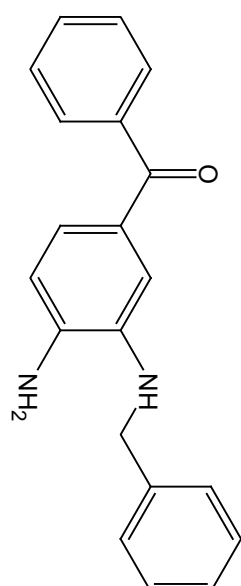
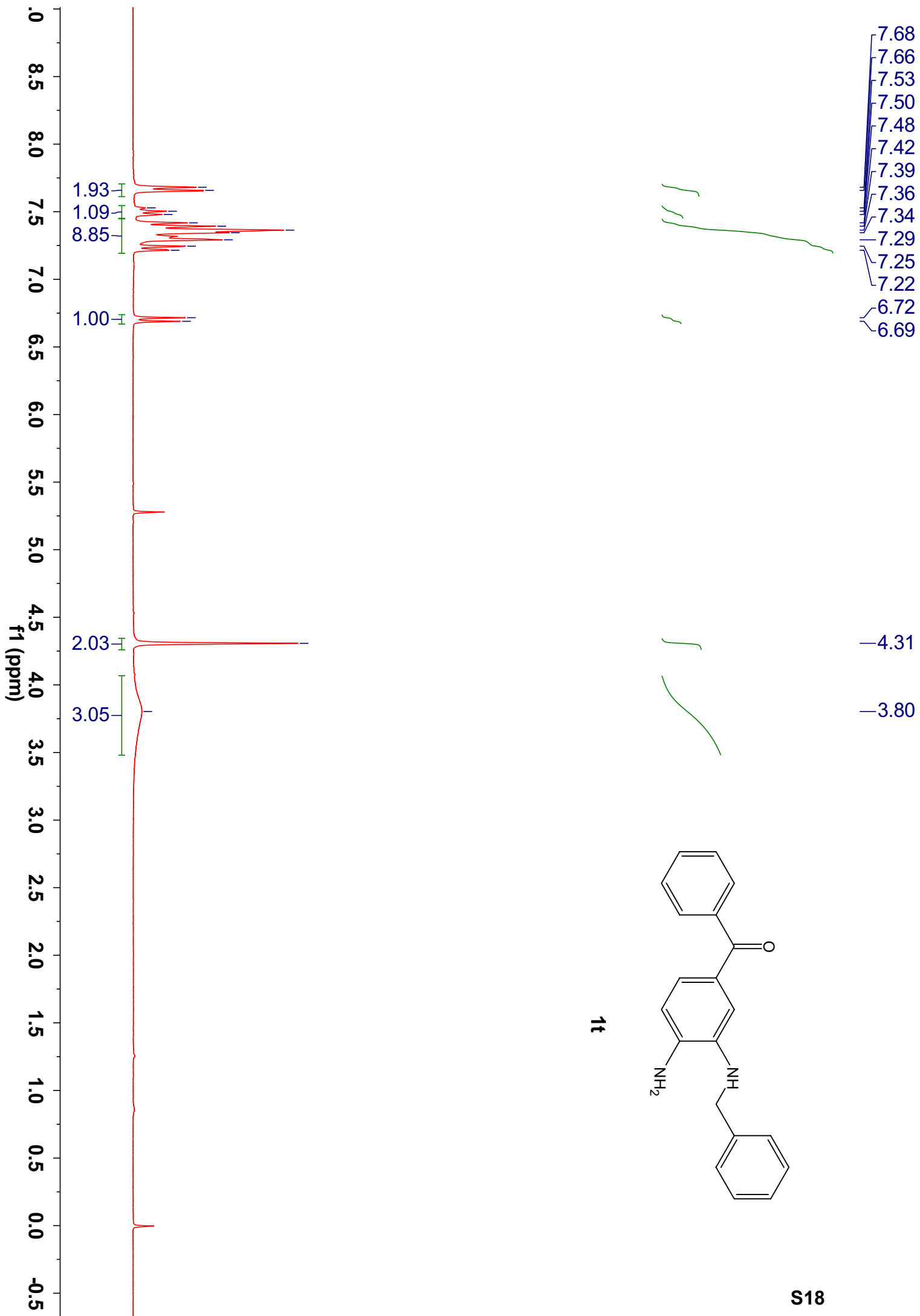
f1 (ppm)

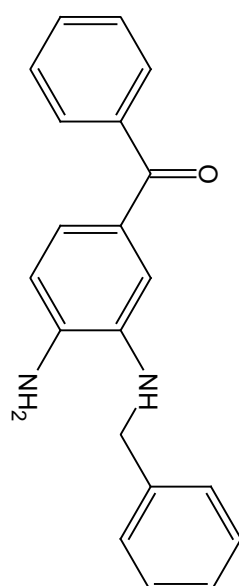
**1r**



1r



**1t**

**1t**

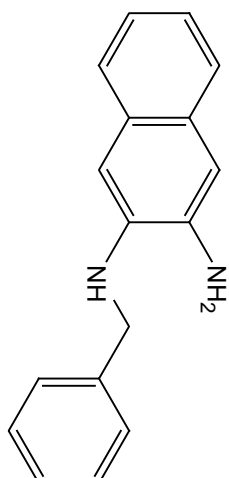
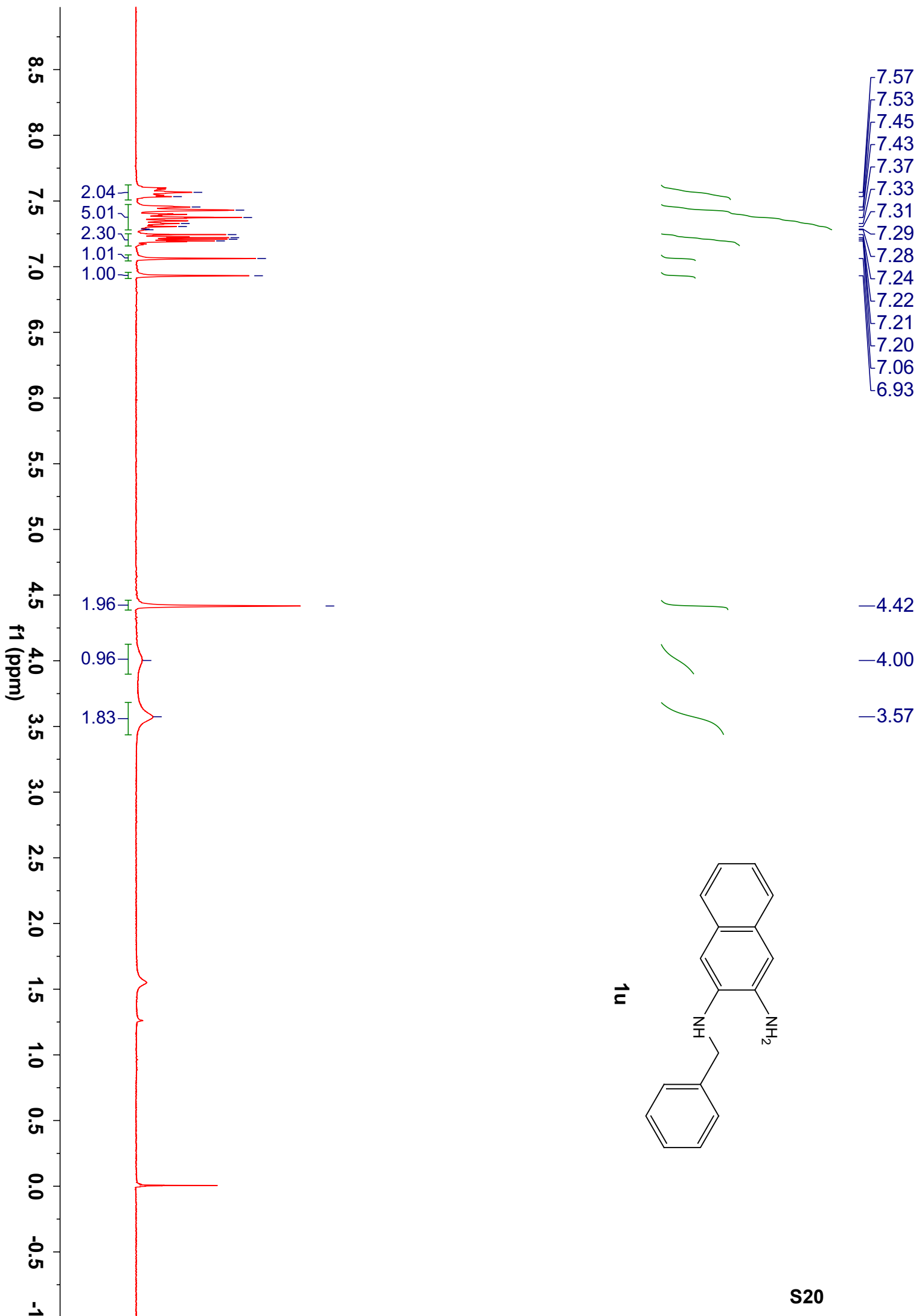
—48.78

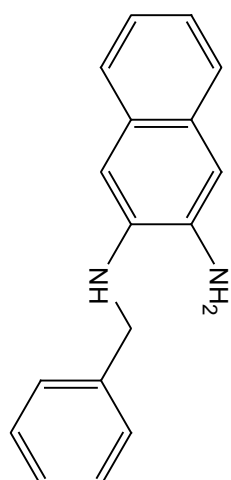
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138.99
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135.98
131.39
129.71
129.05
128.78
128.08
128.04
127.54
124.48
114.44
114.19

—196.03

f1 (ppm)

10 200 190 180 170 160 150 140 130 120 110 100 90 80 70 60 50 40 30 20 10 0

**1u**

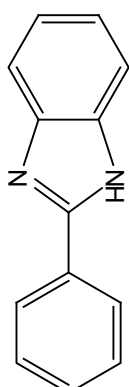
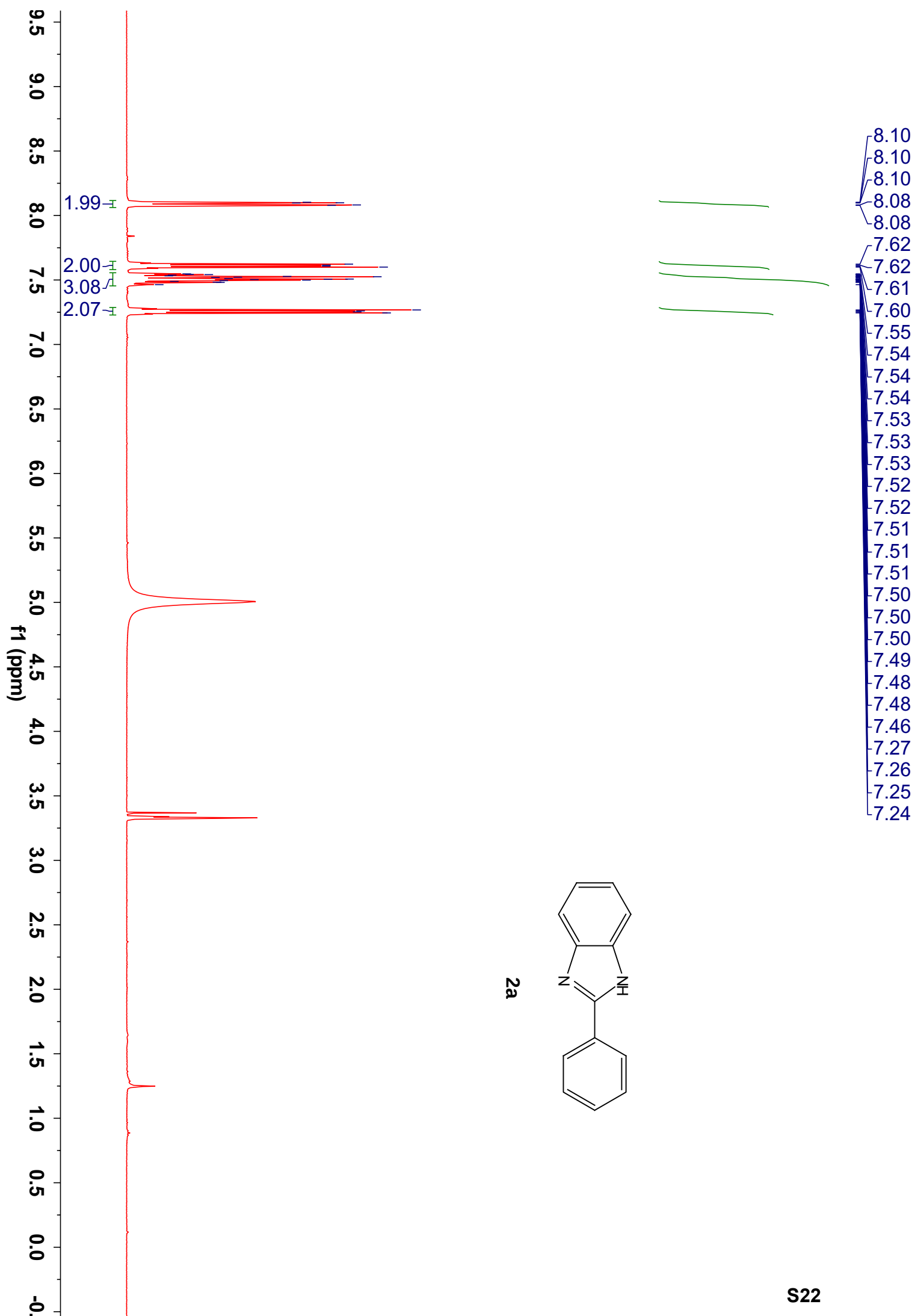


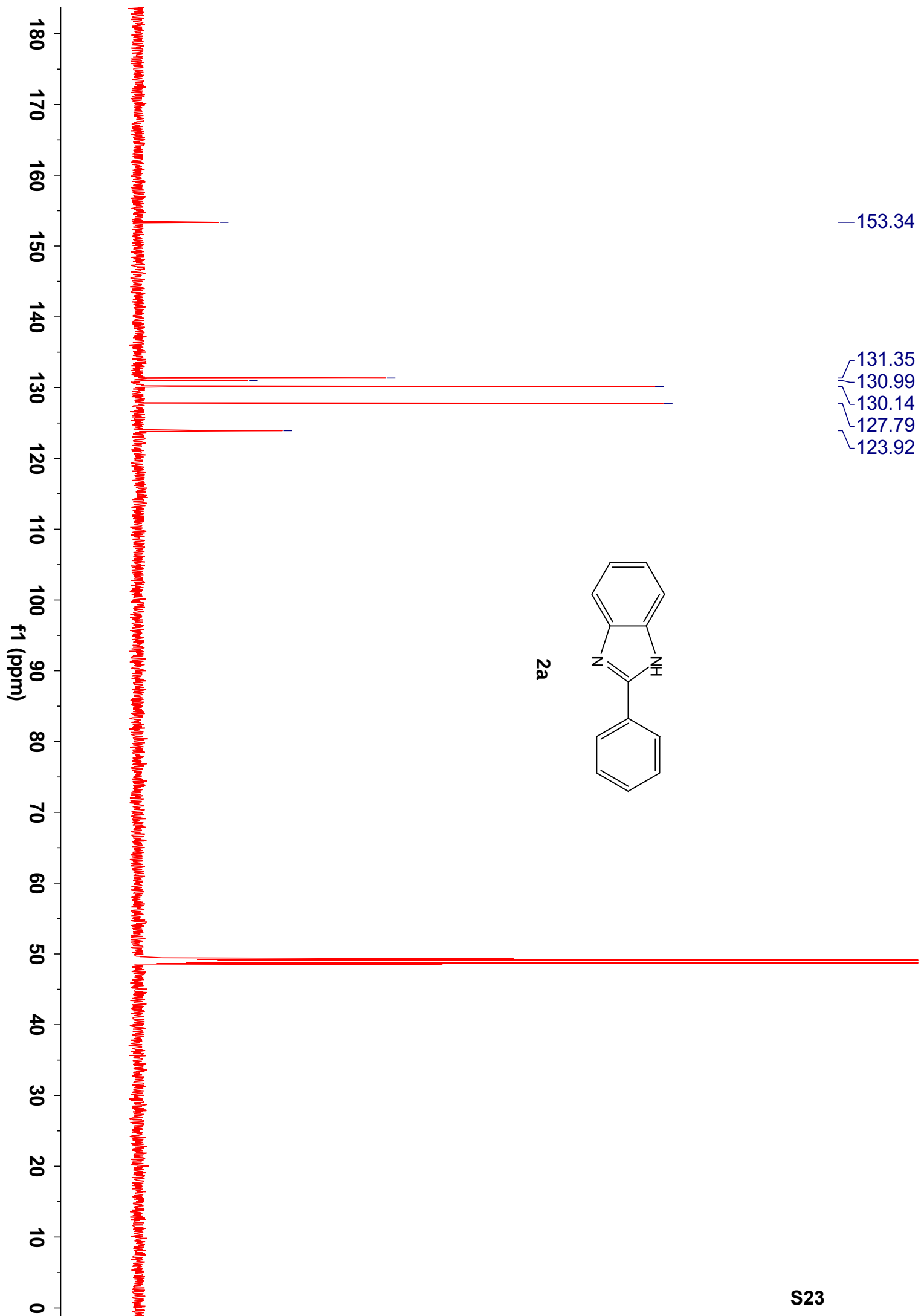
1u

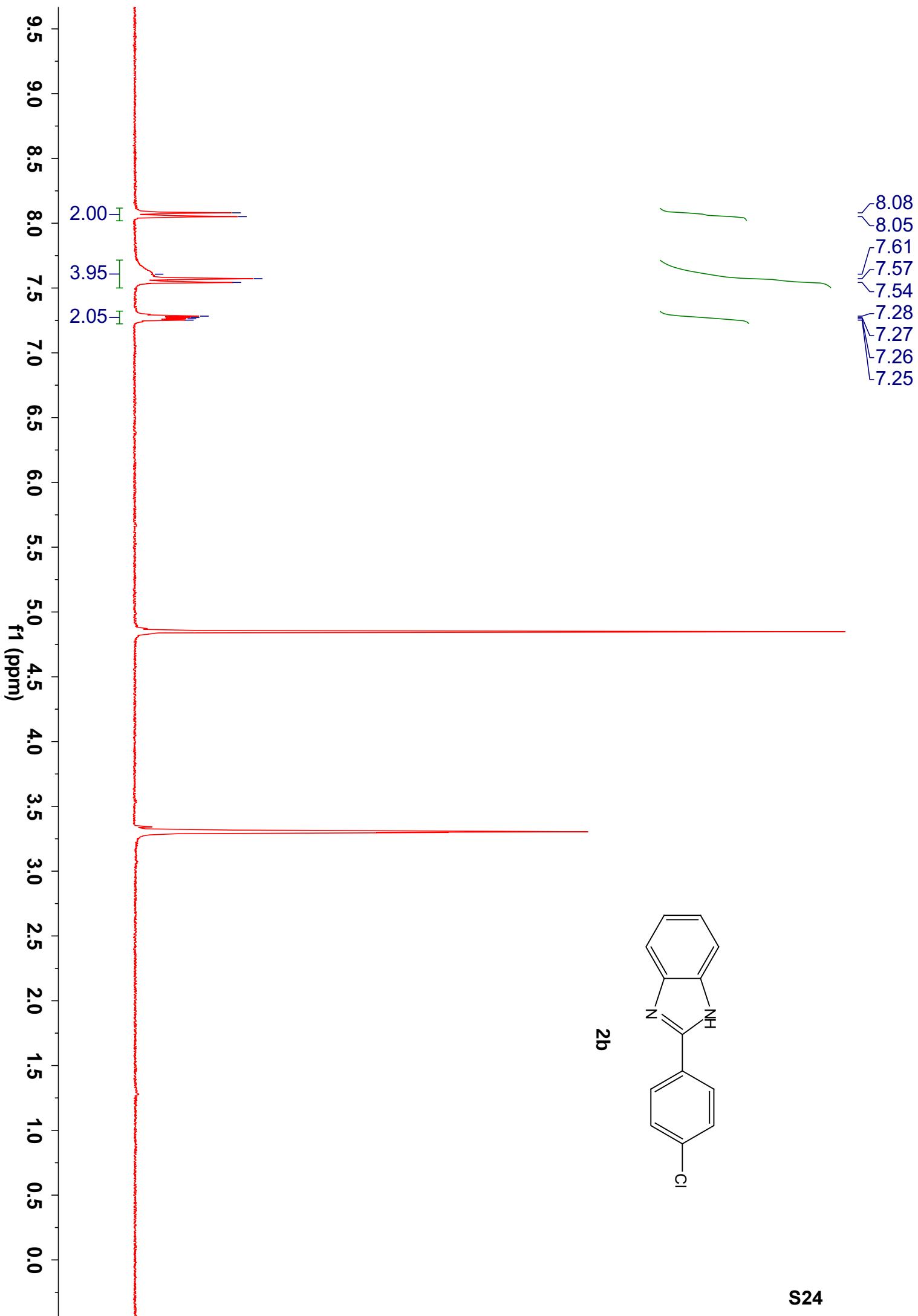
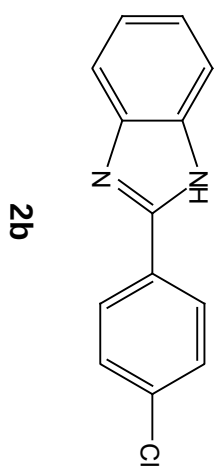
139.07
138.87
135.67
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128.82
128.68
128.06
127.55
126.03
125.63
123.48
122.95
111.52
106.30

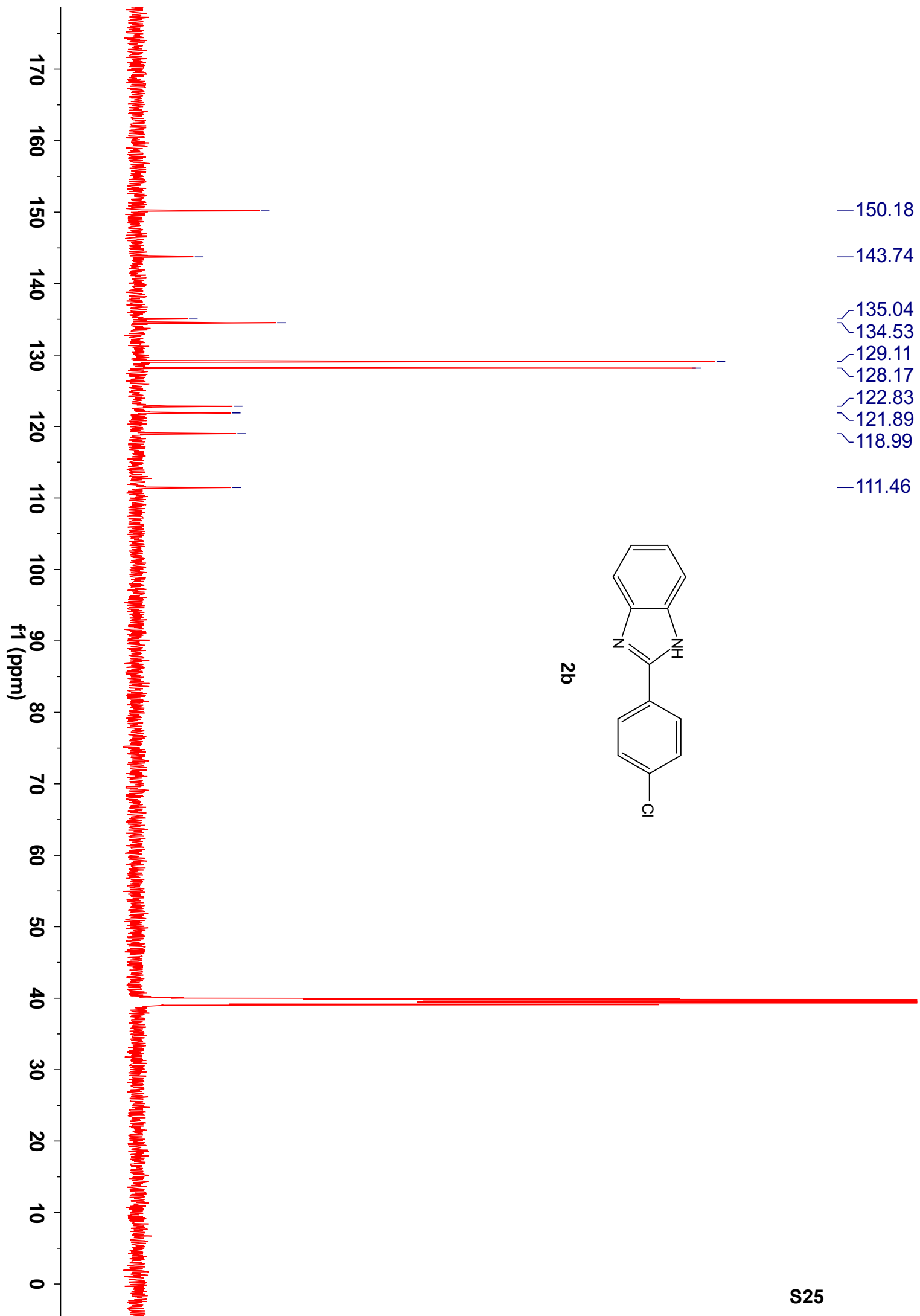
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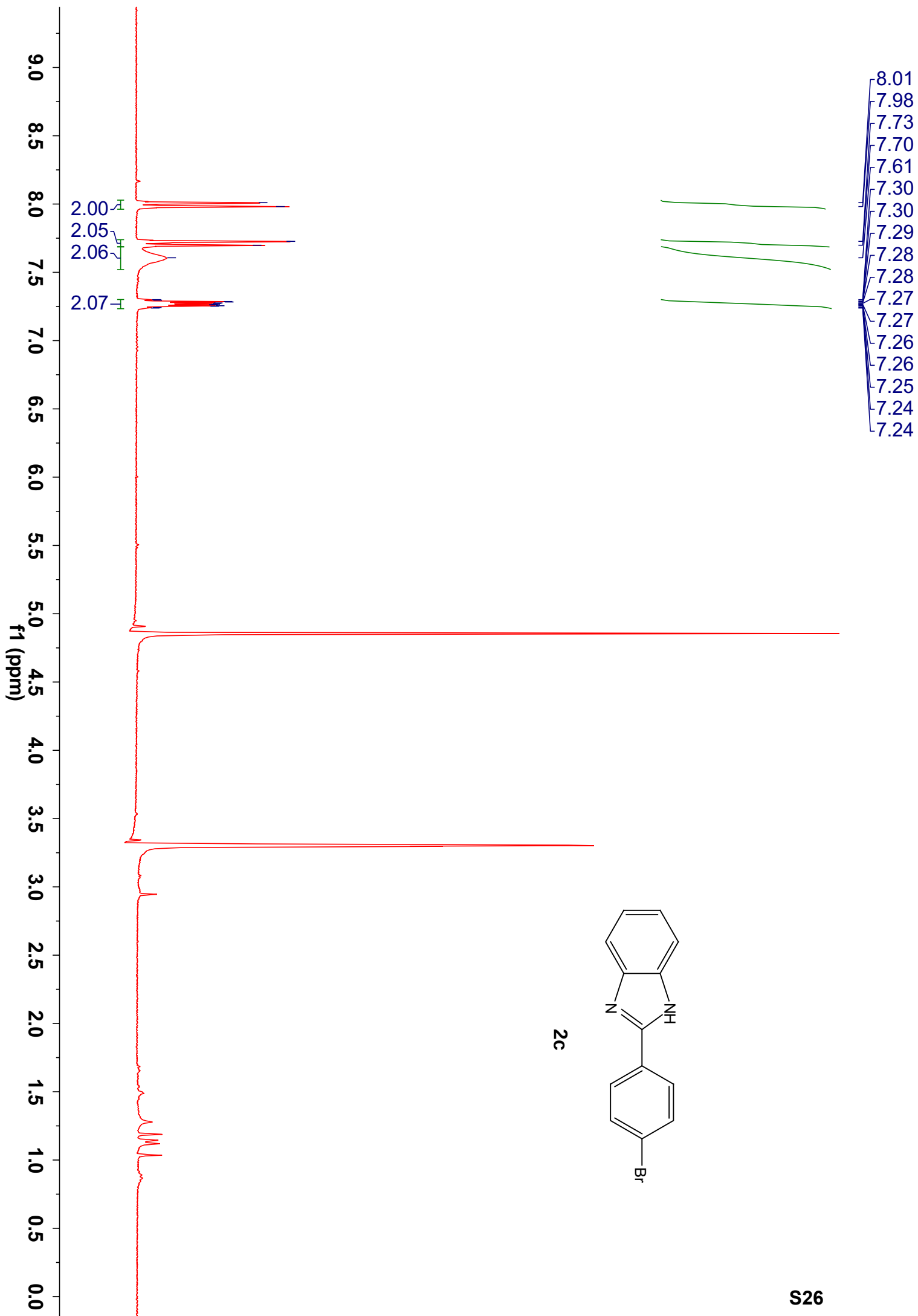
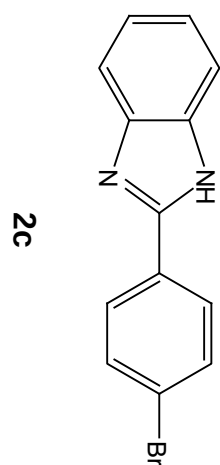
f1 (ppm)

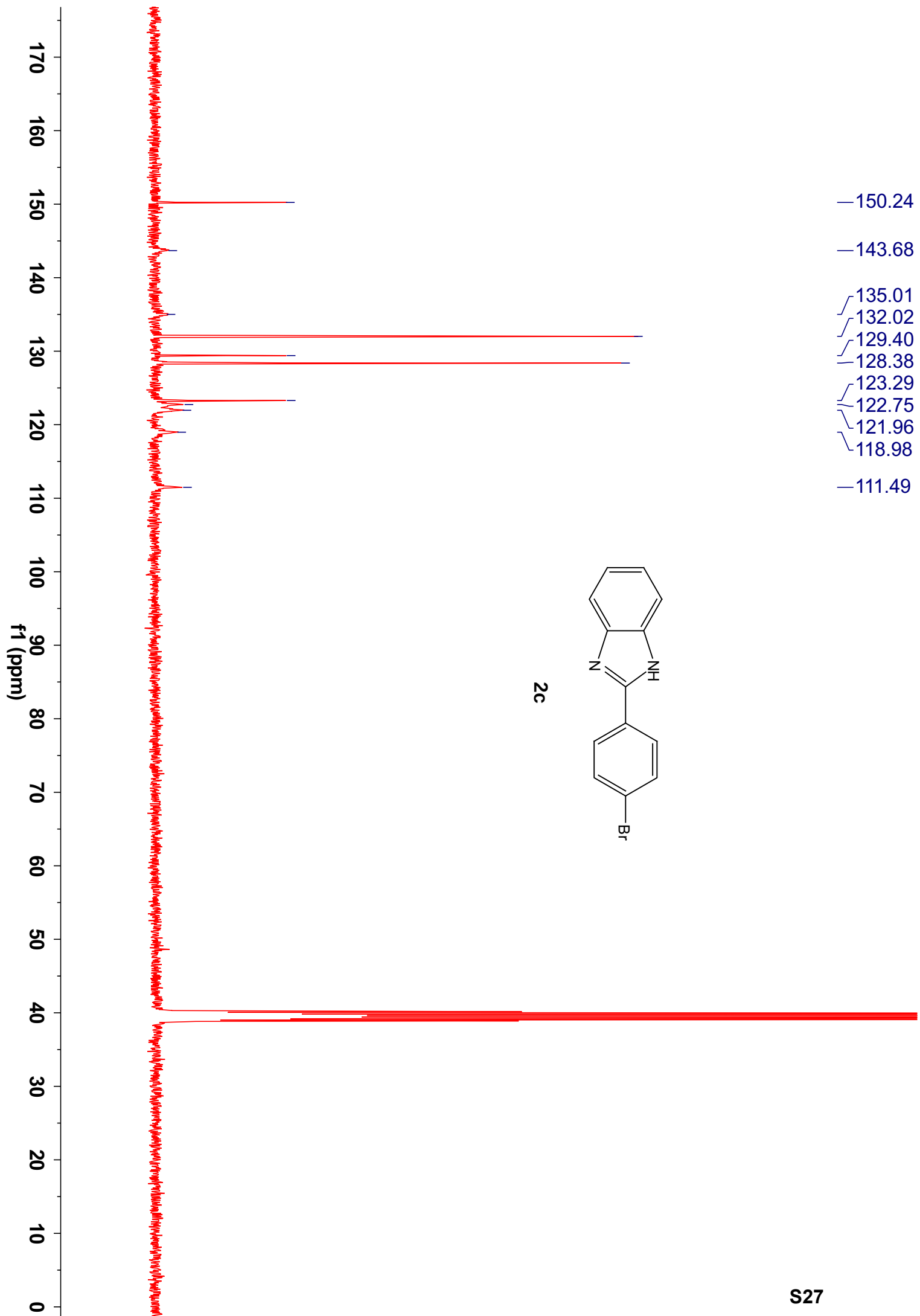
**2a**

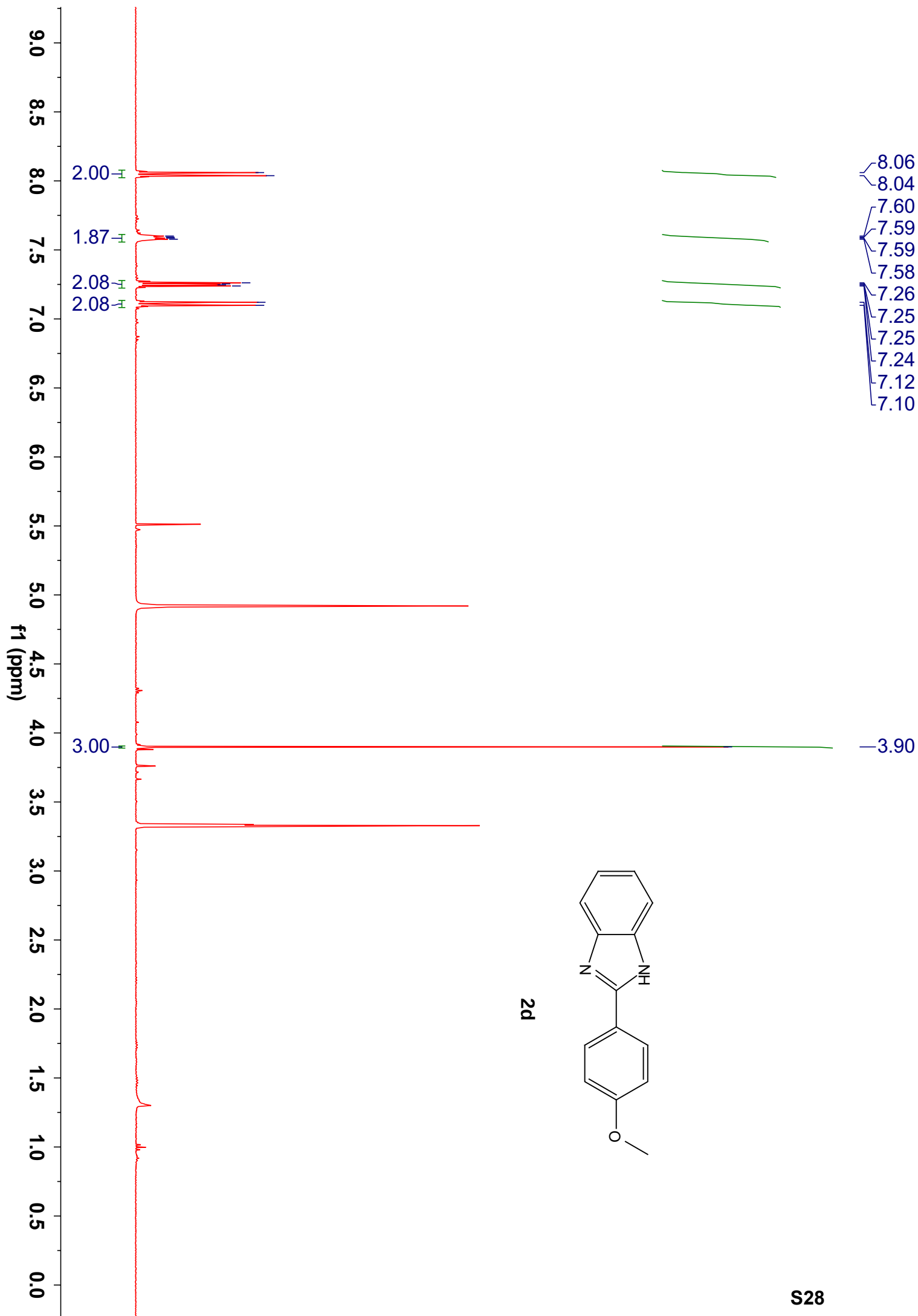


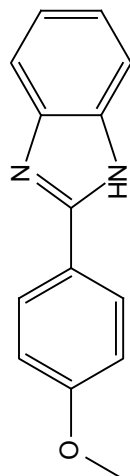




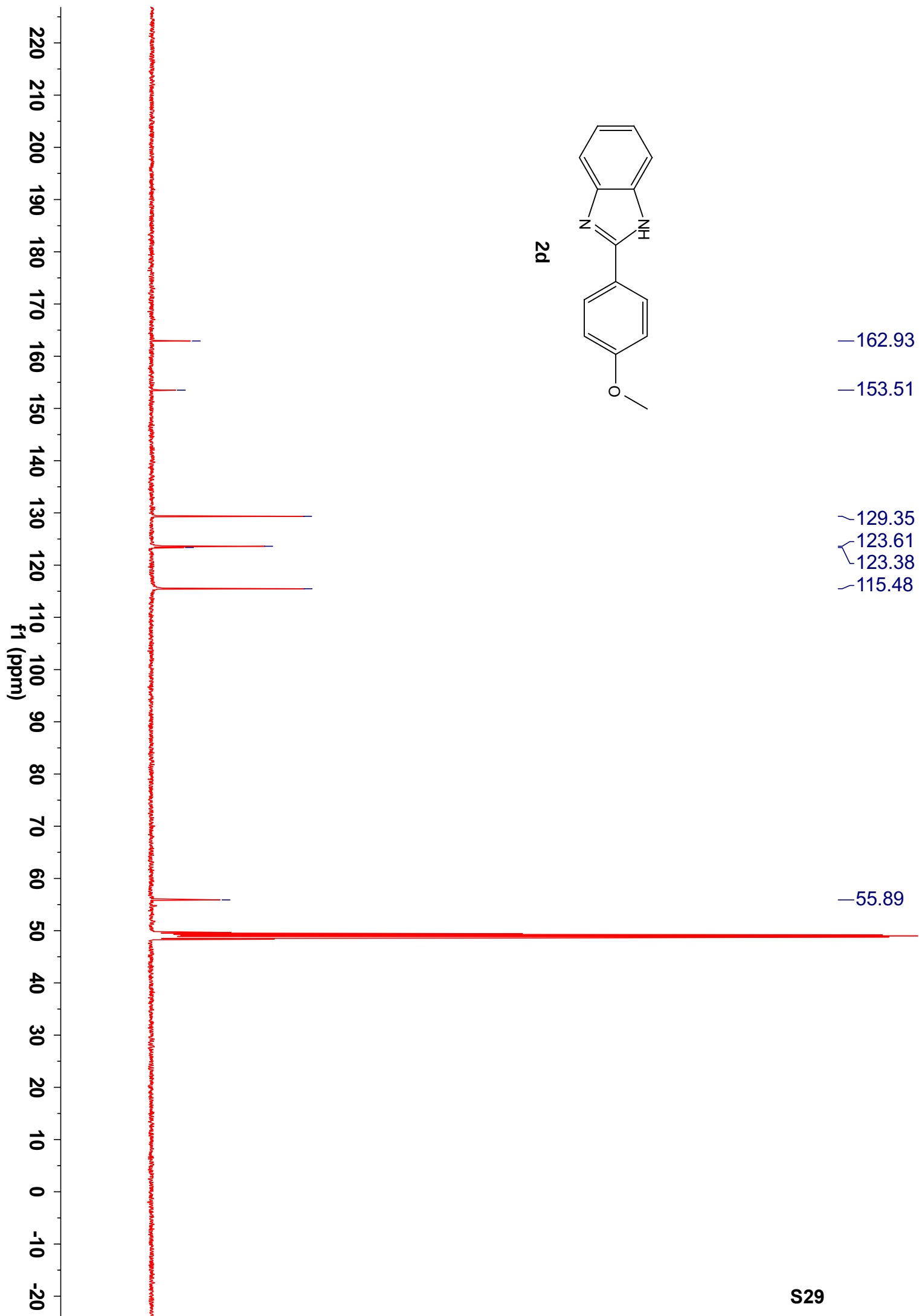


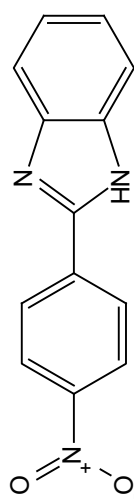
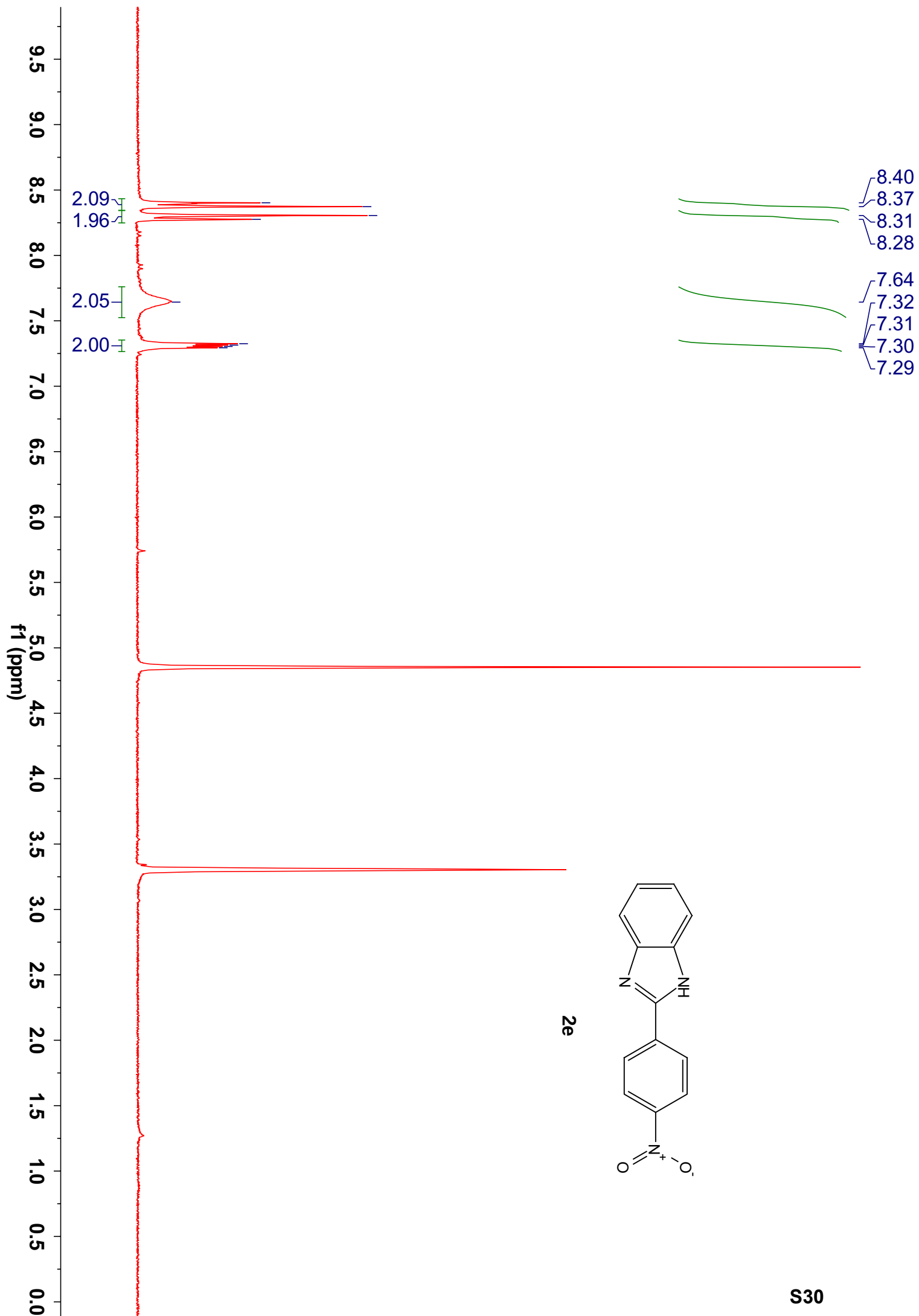


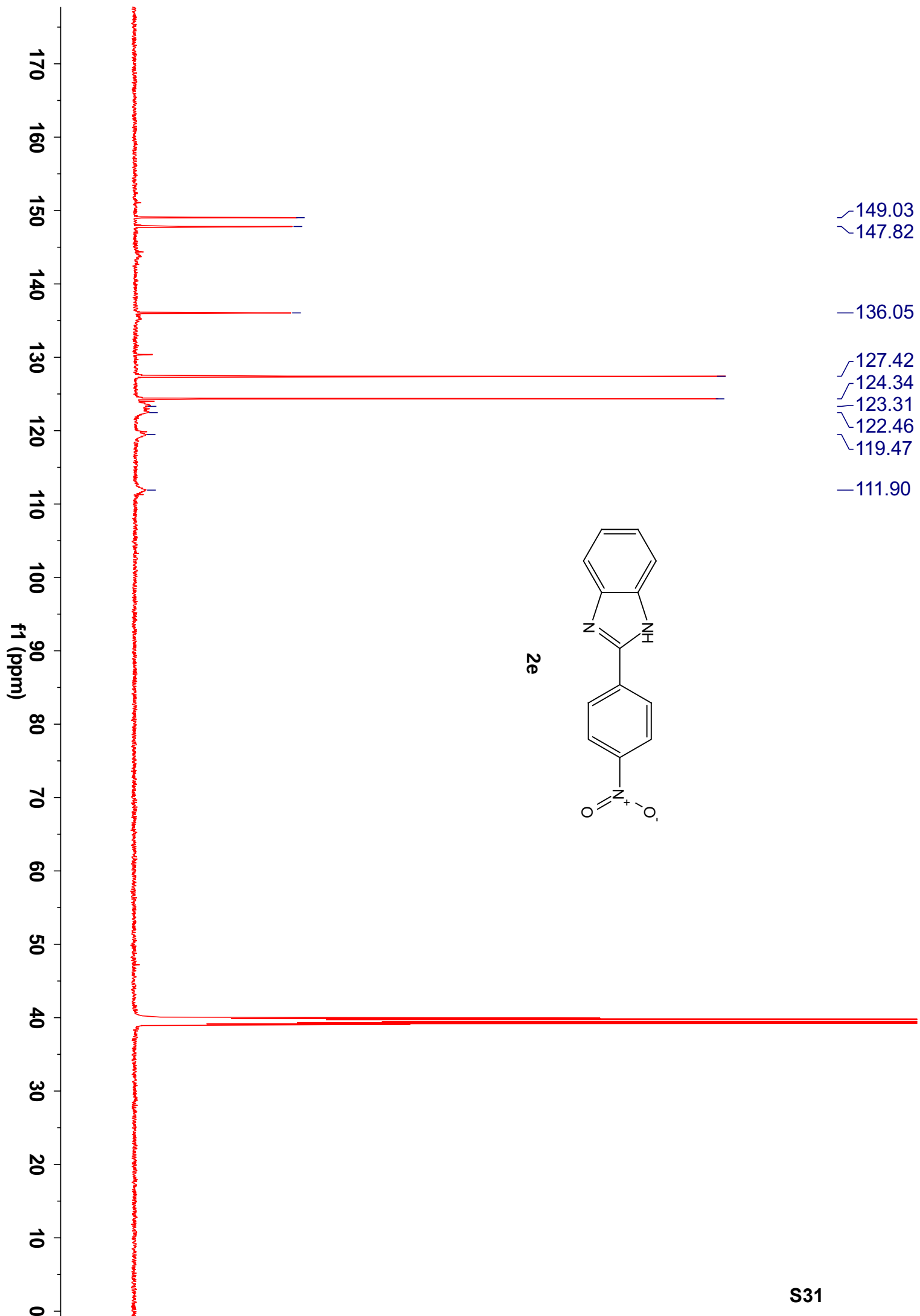


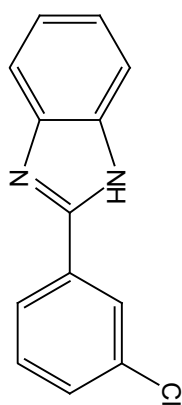


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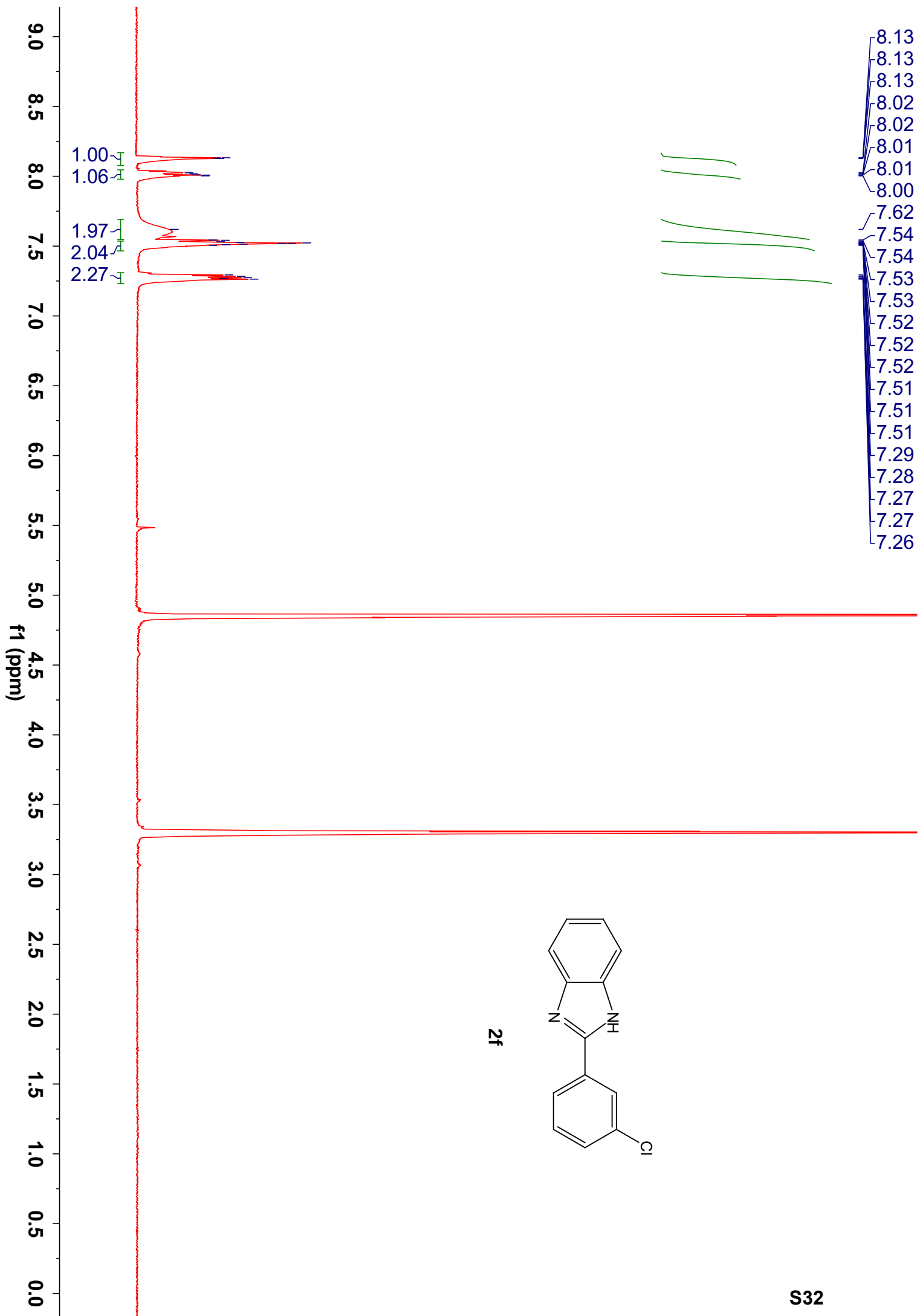


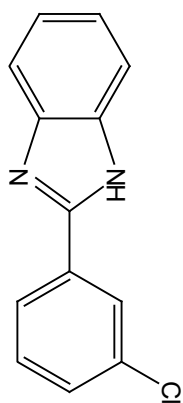
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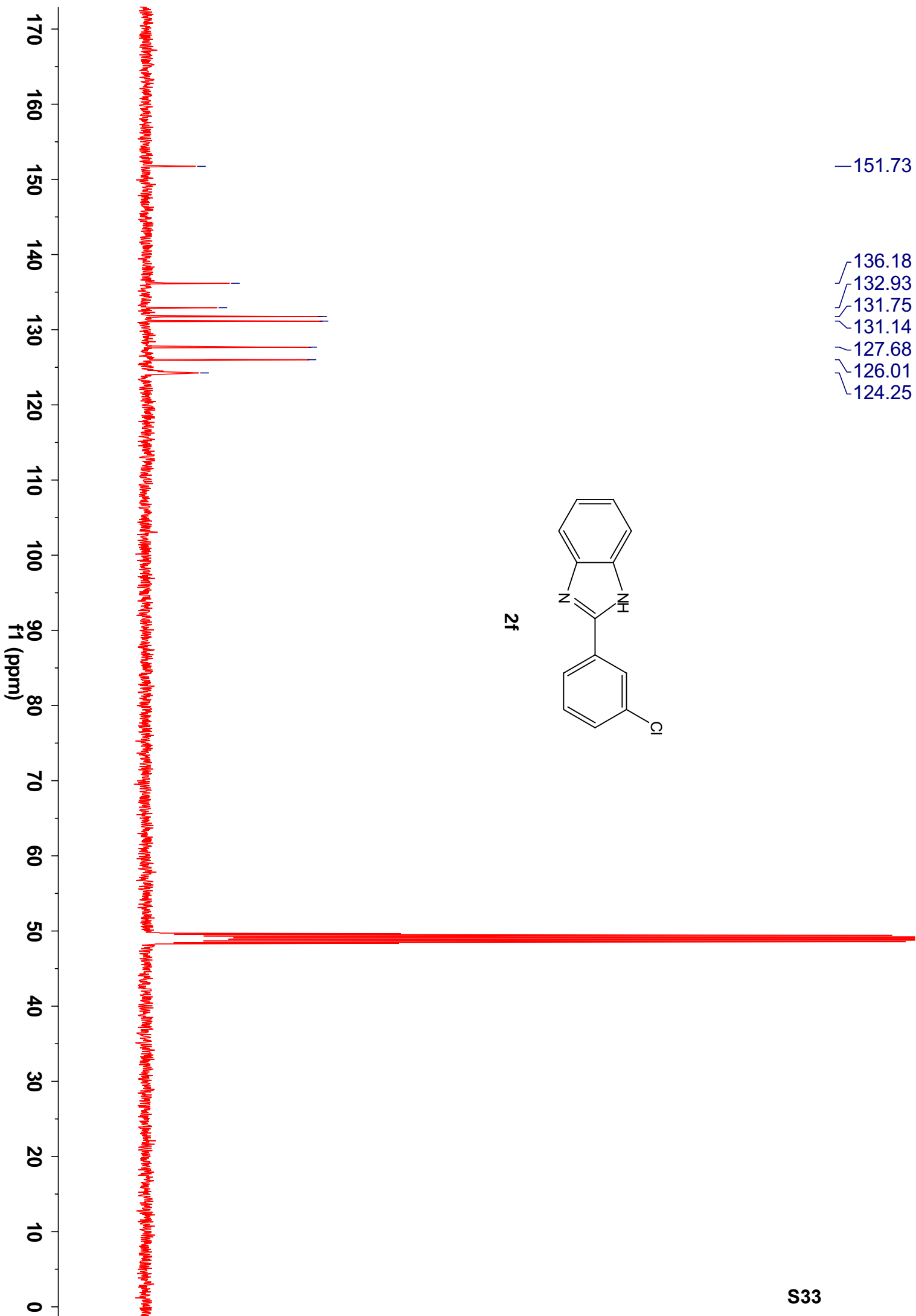


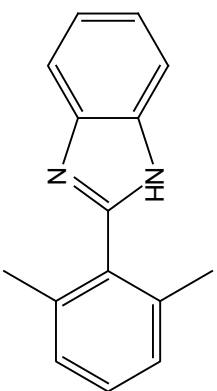
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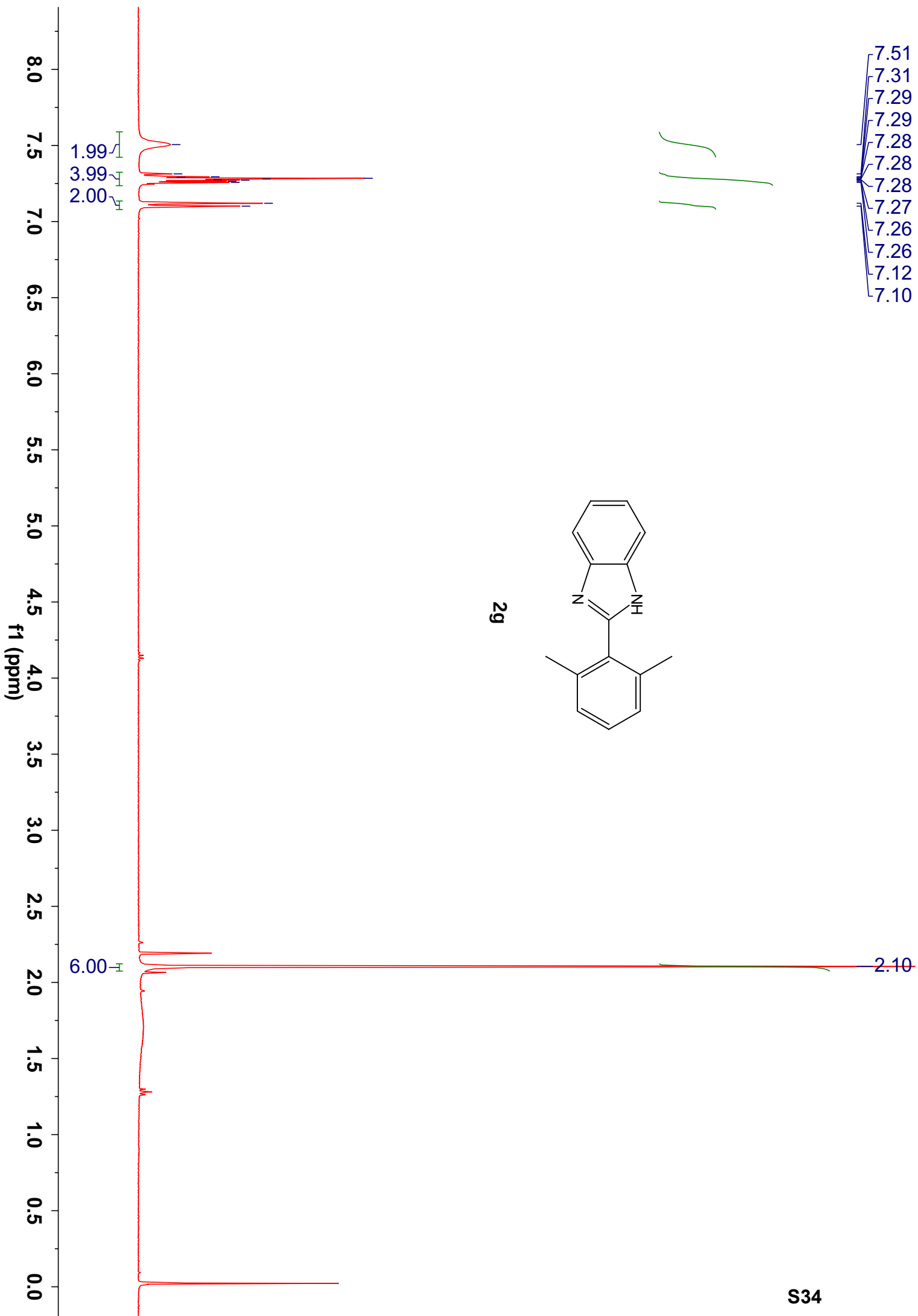


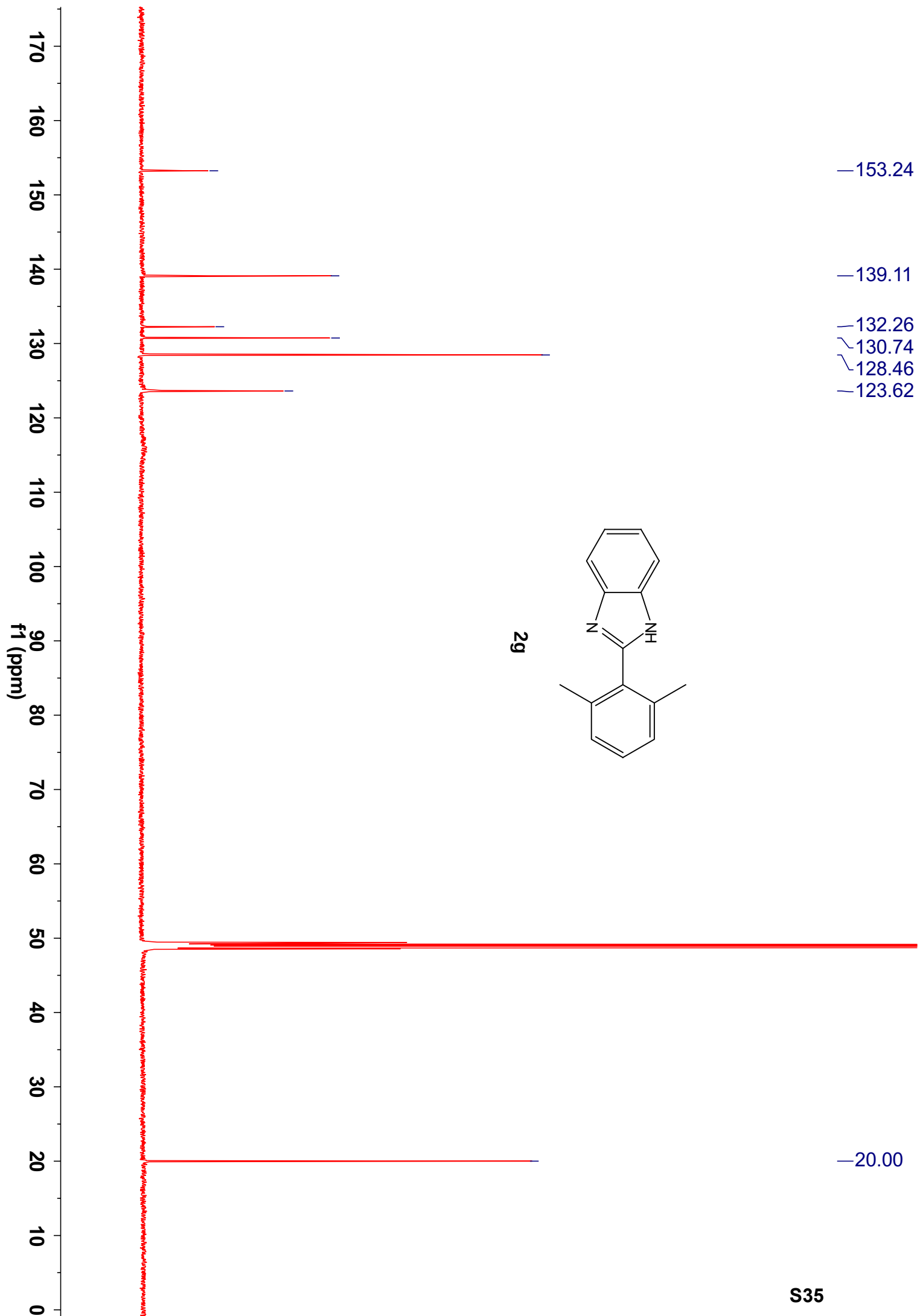
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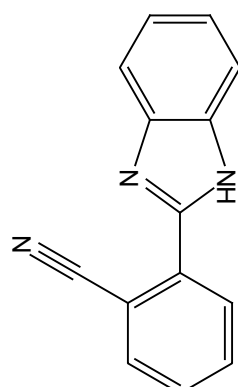




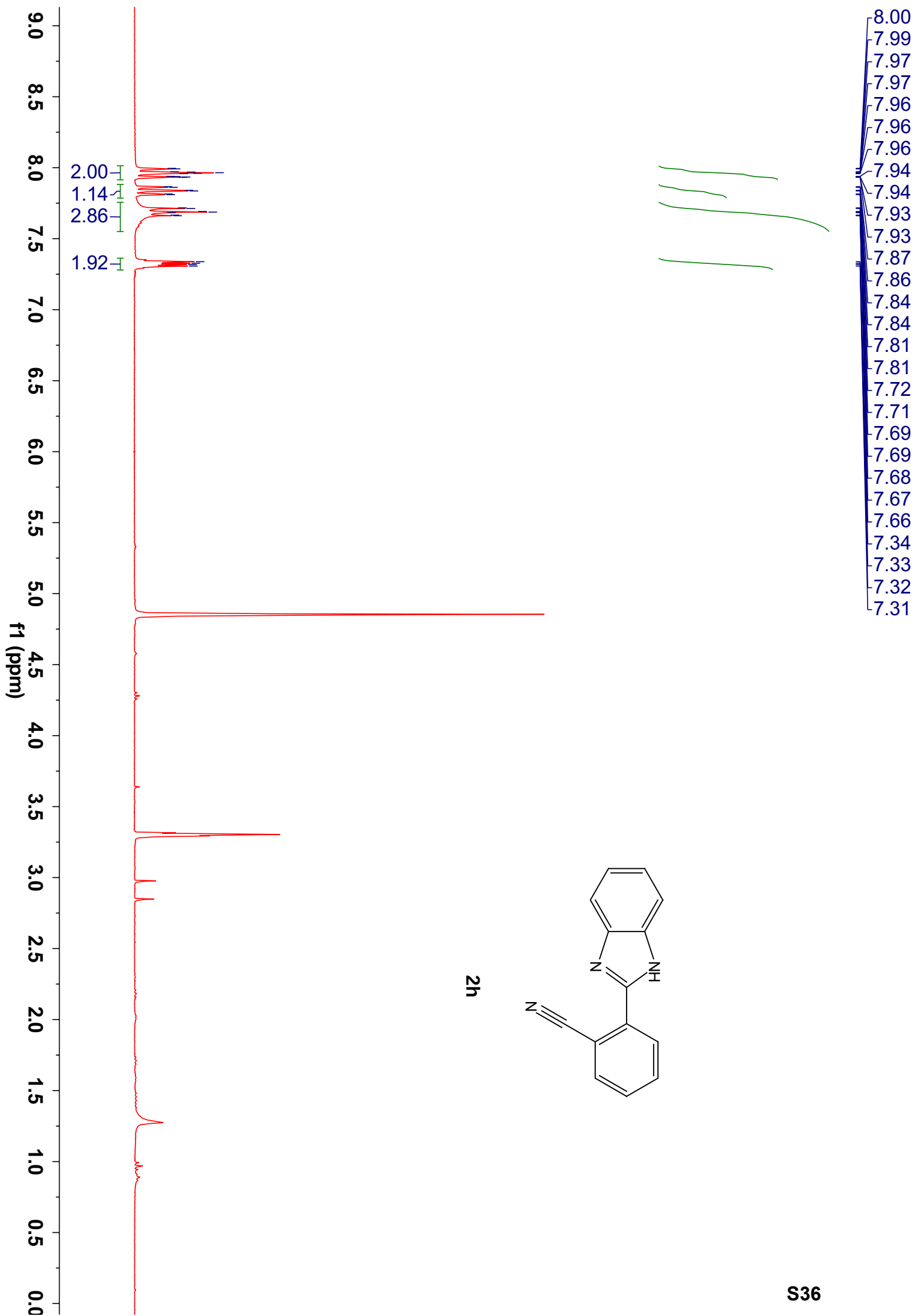
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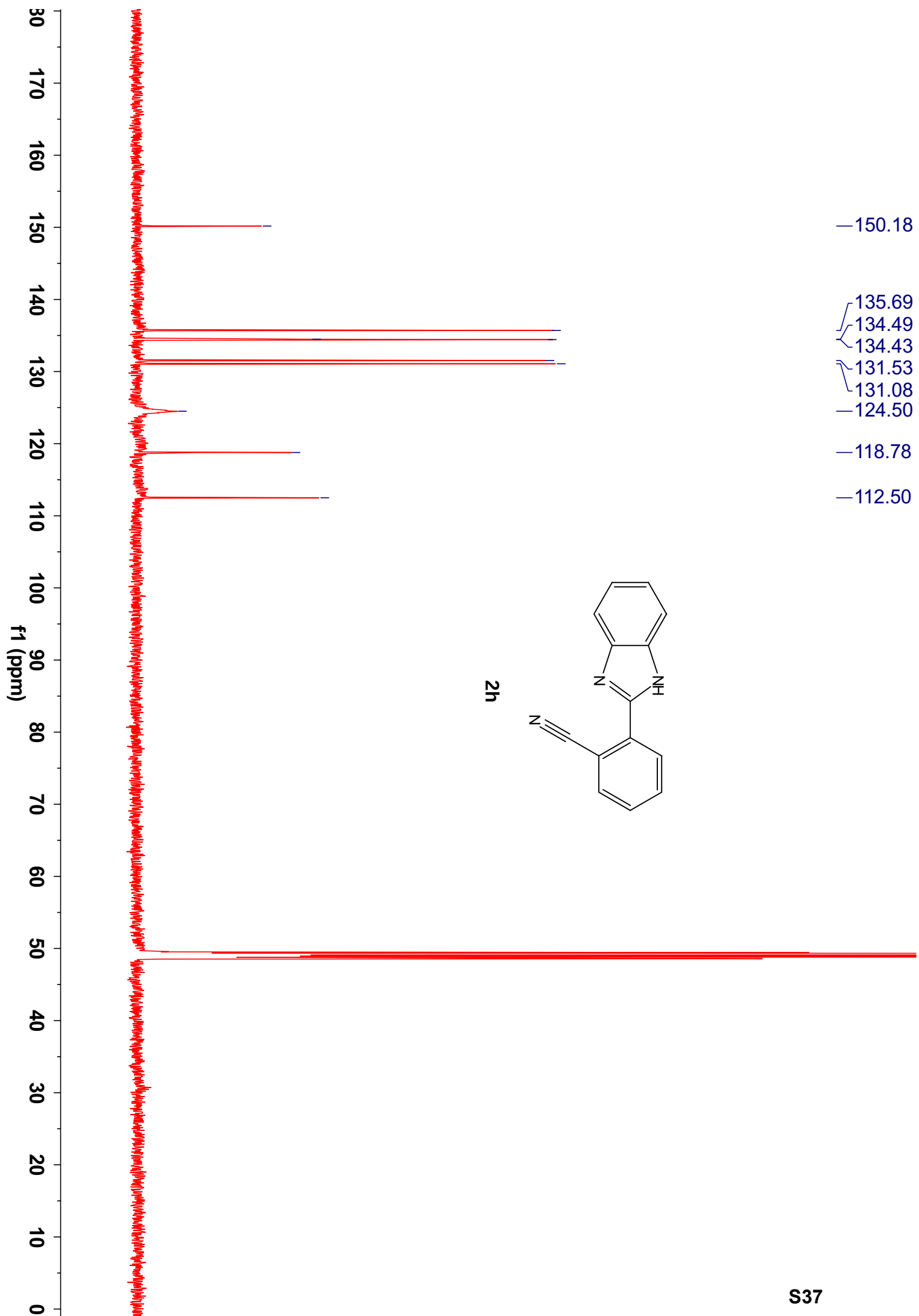


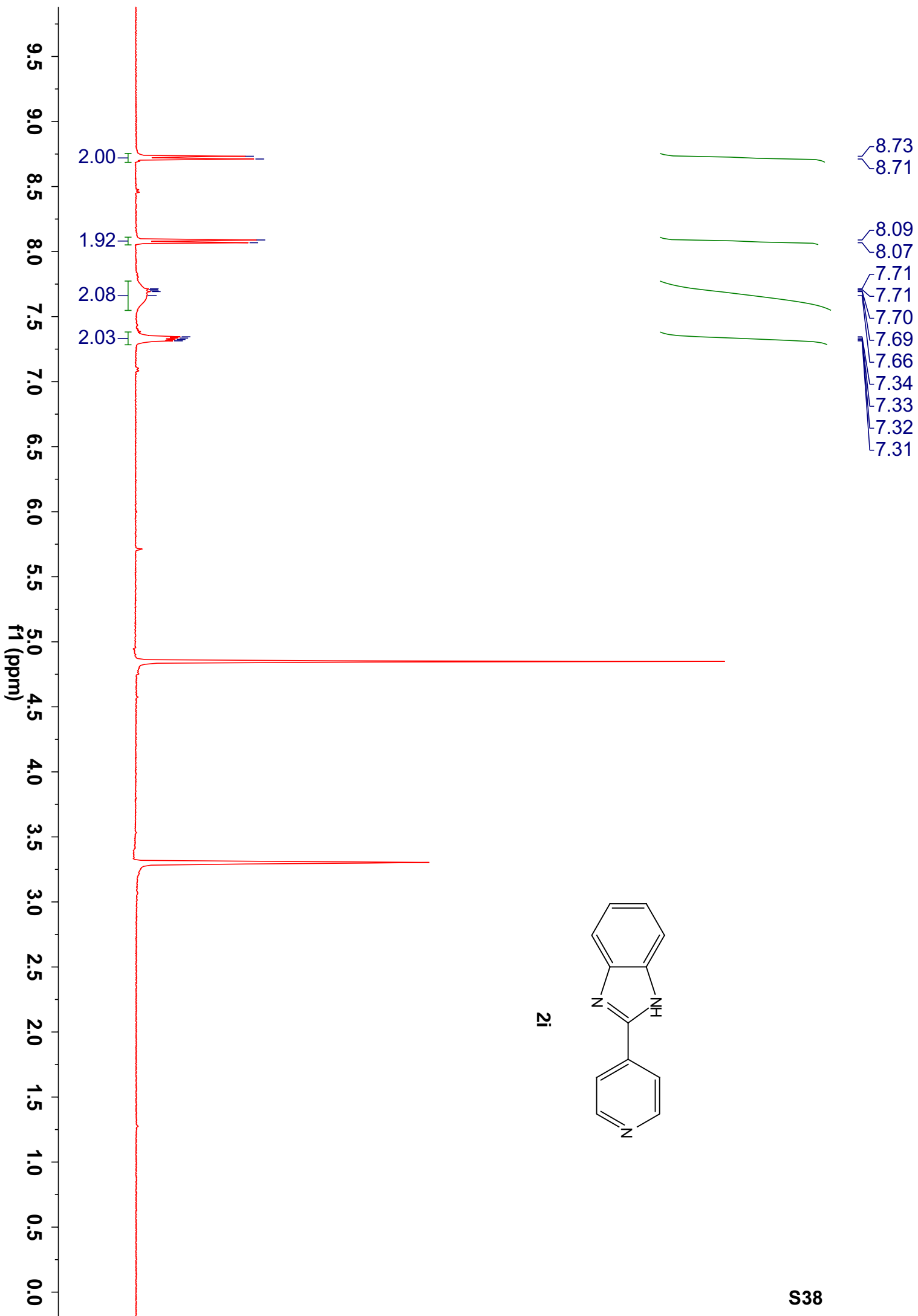
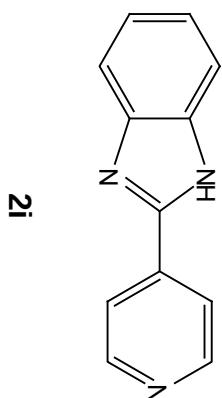


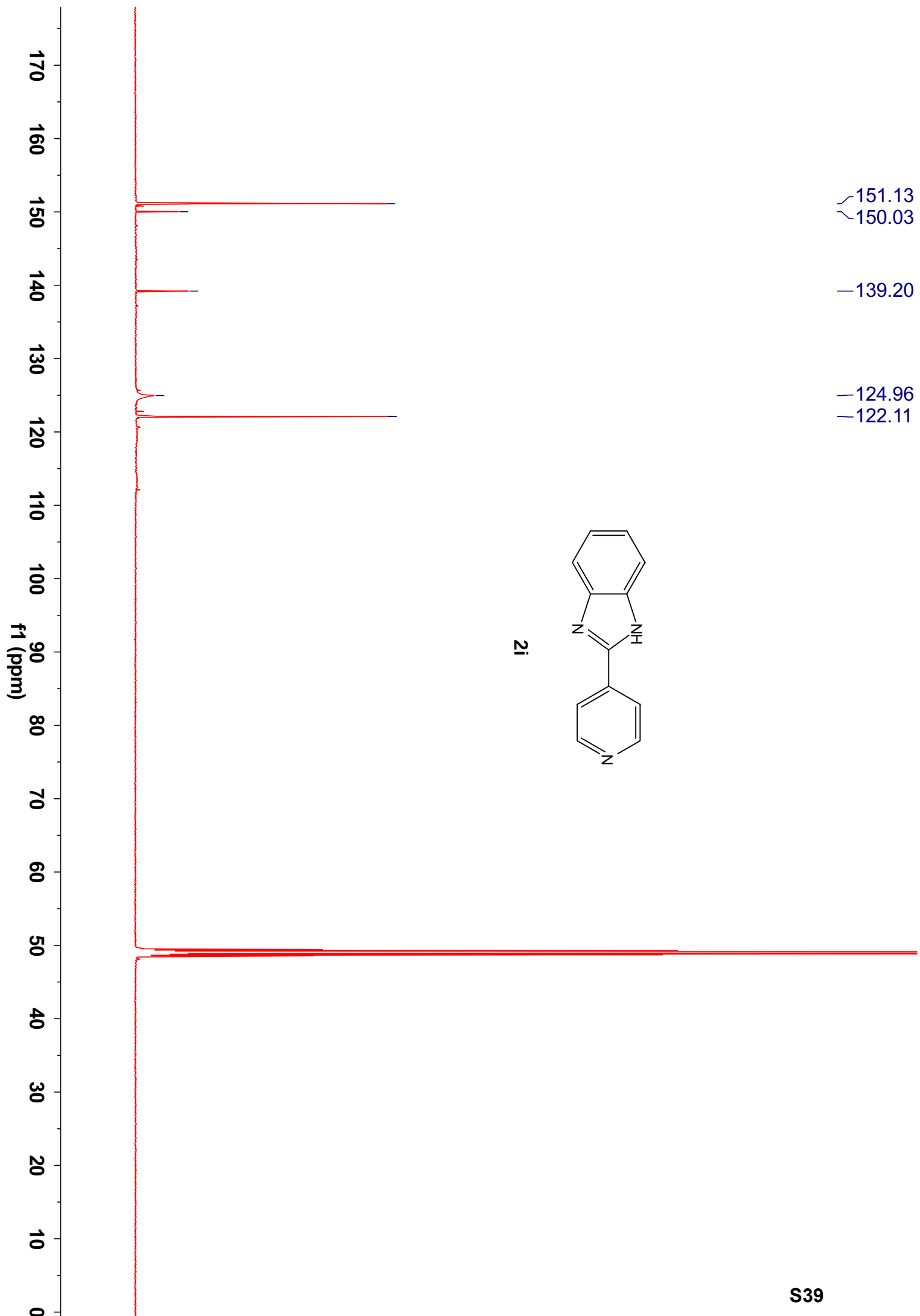
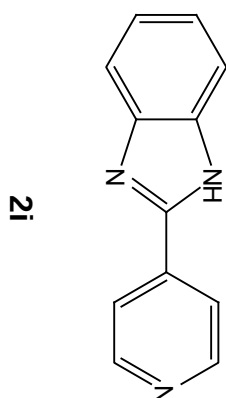


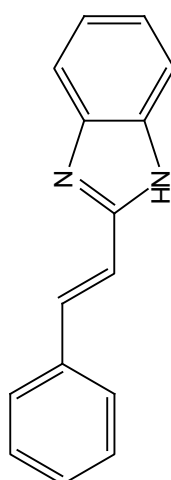
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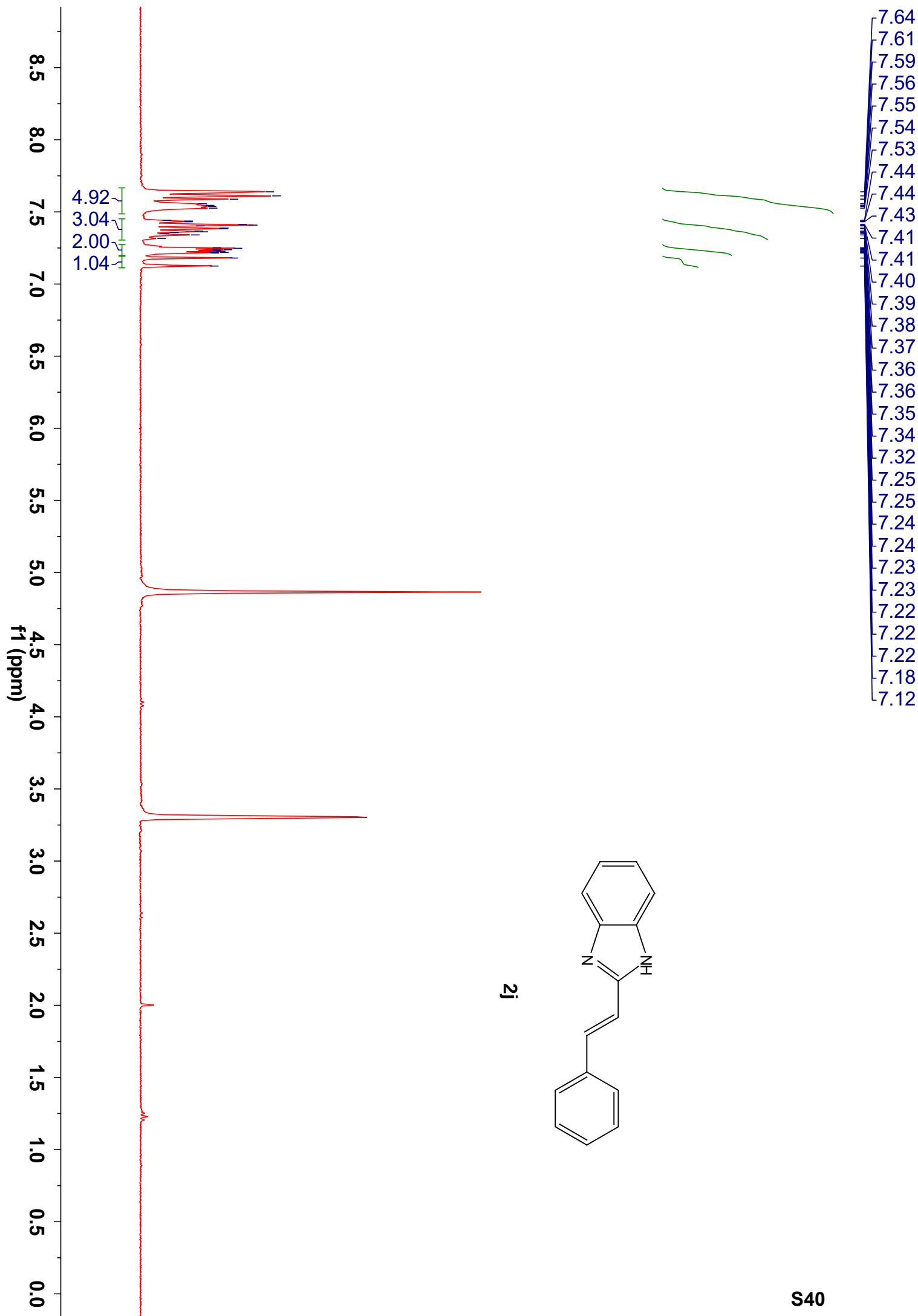


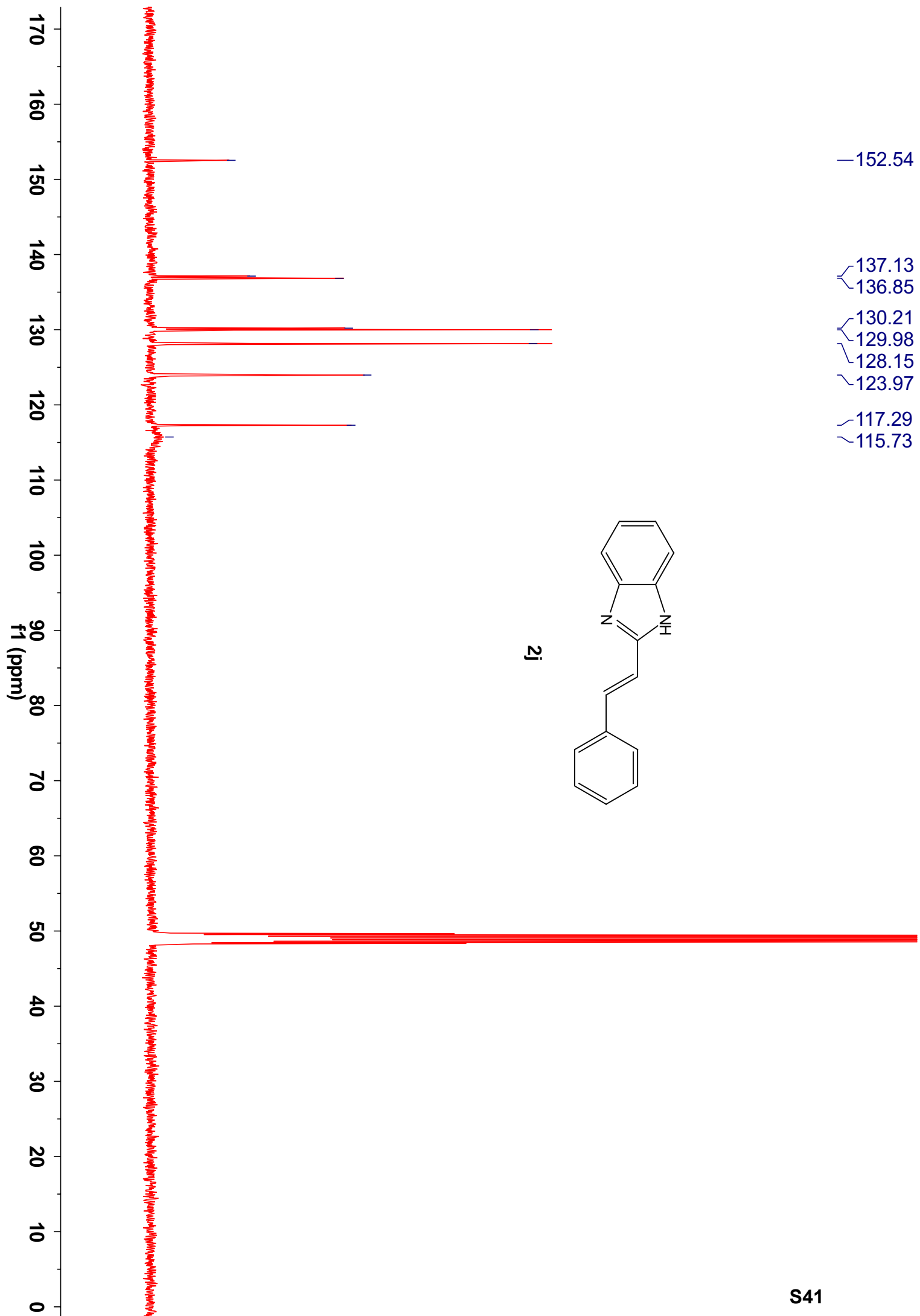


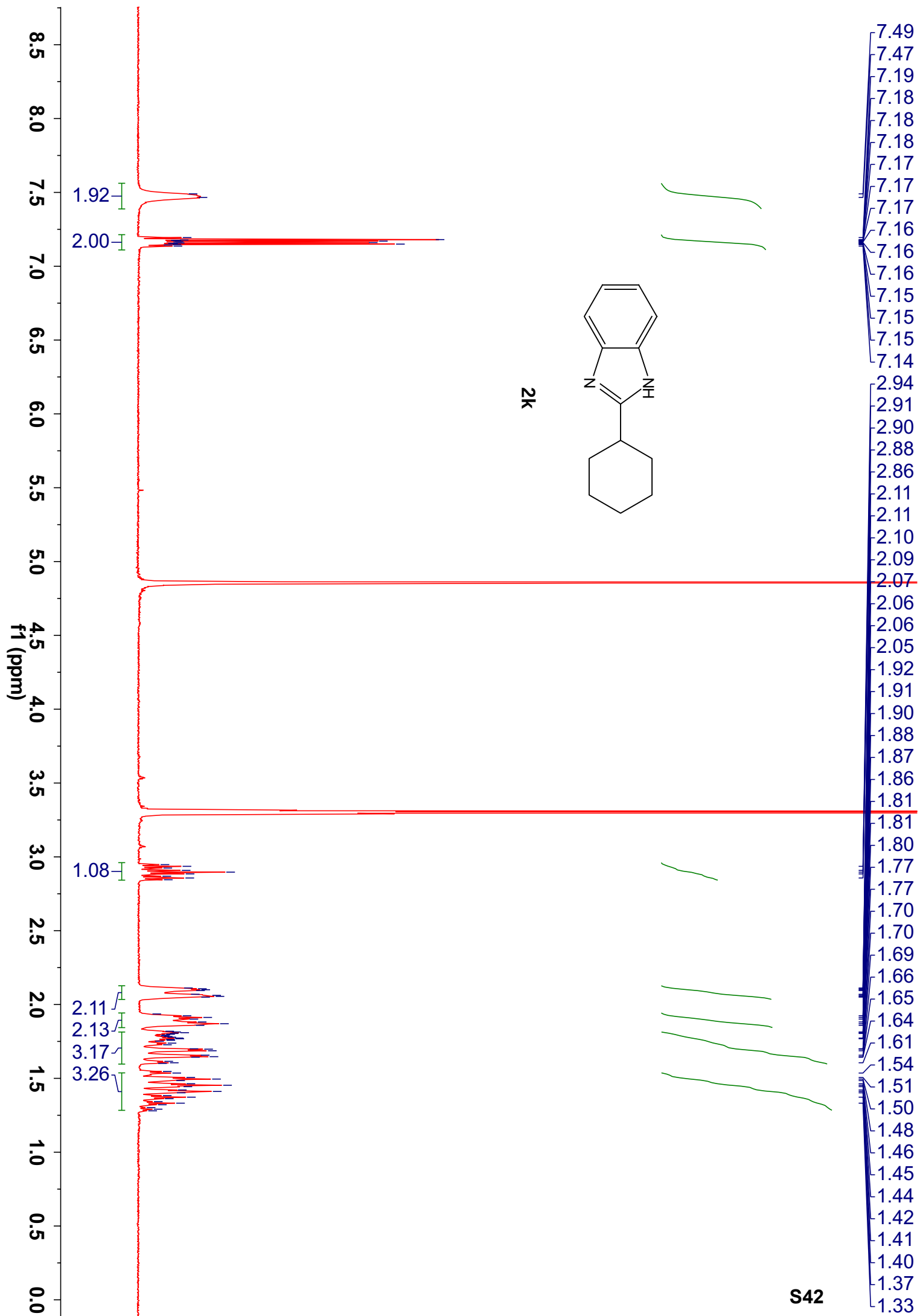


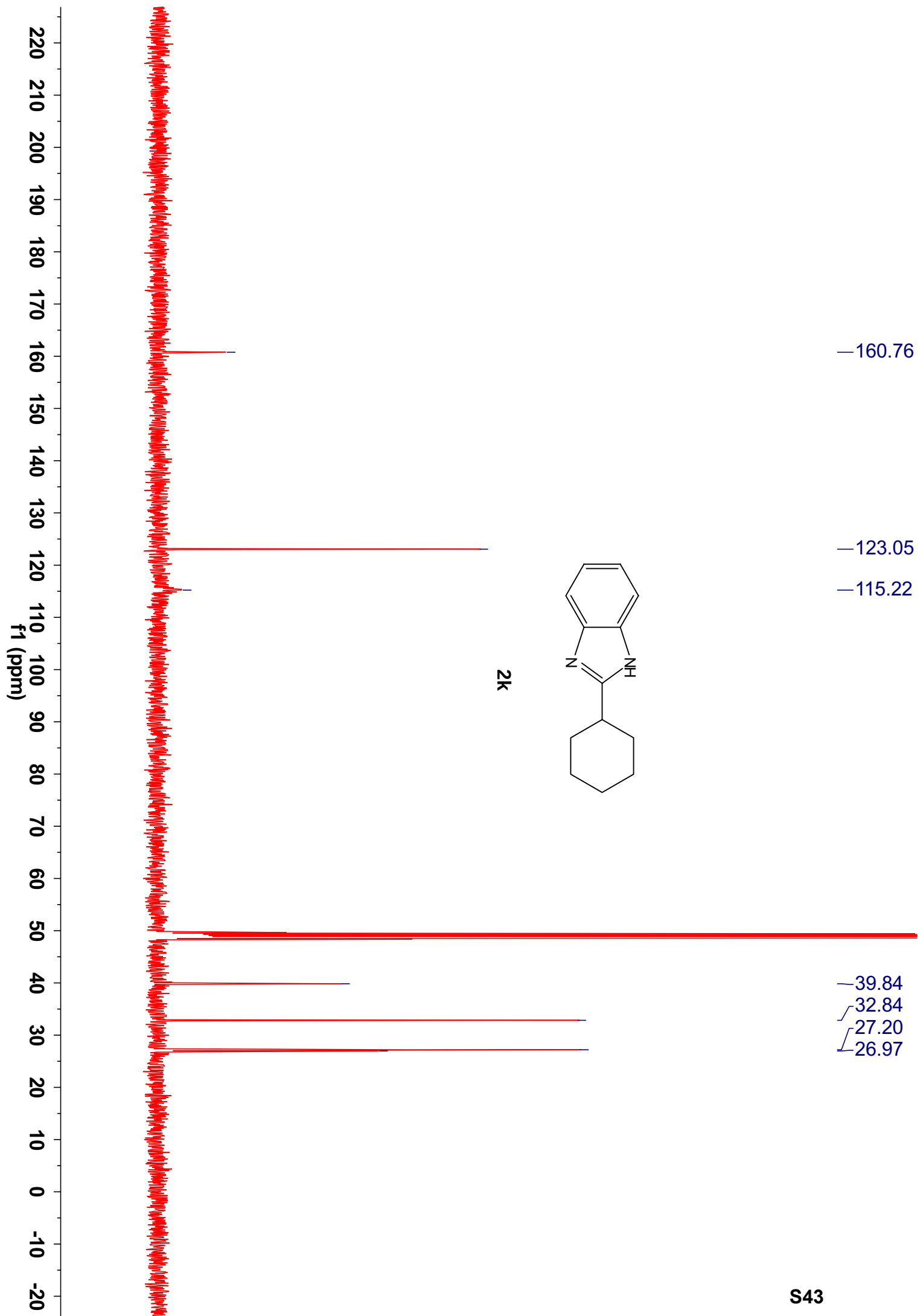


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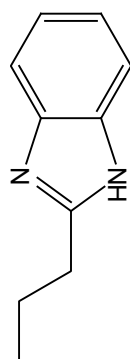




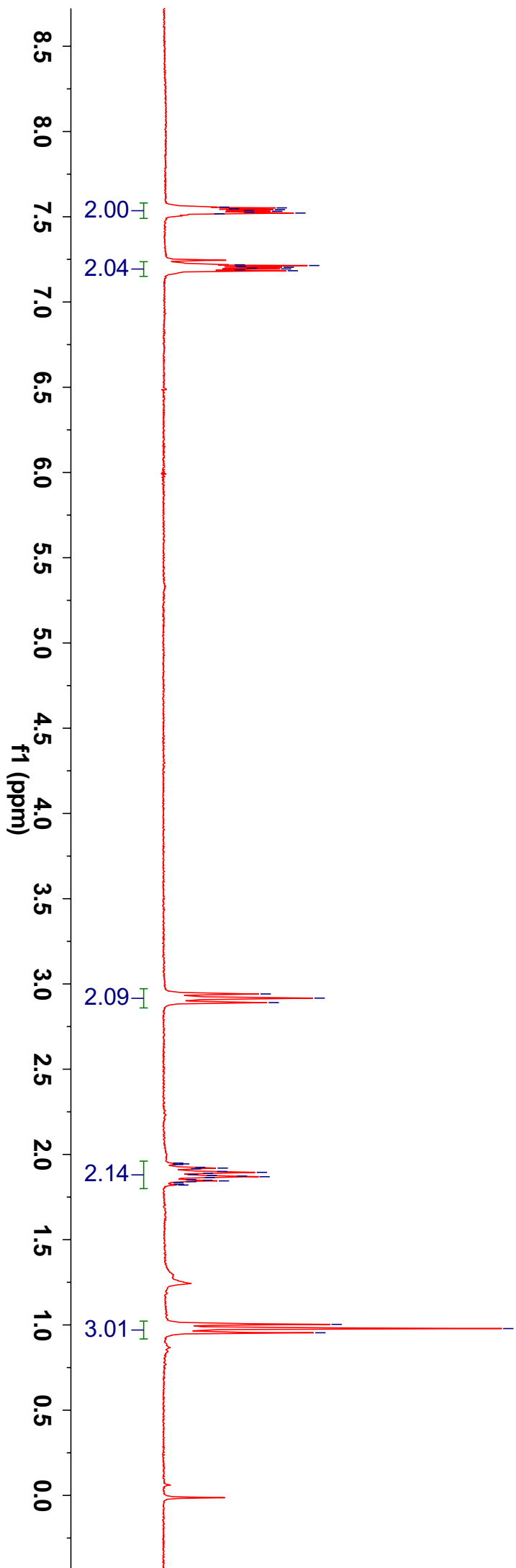


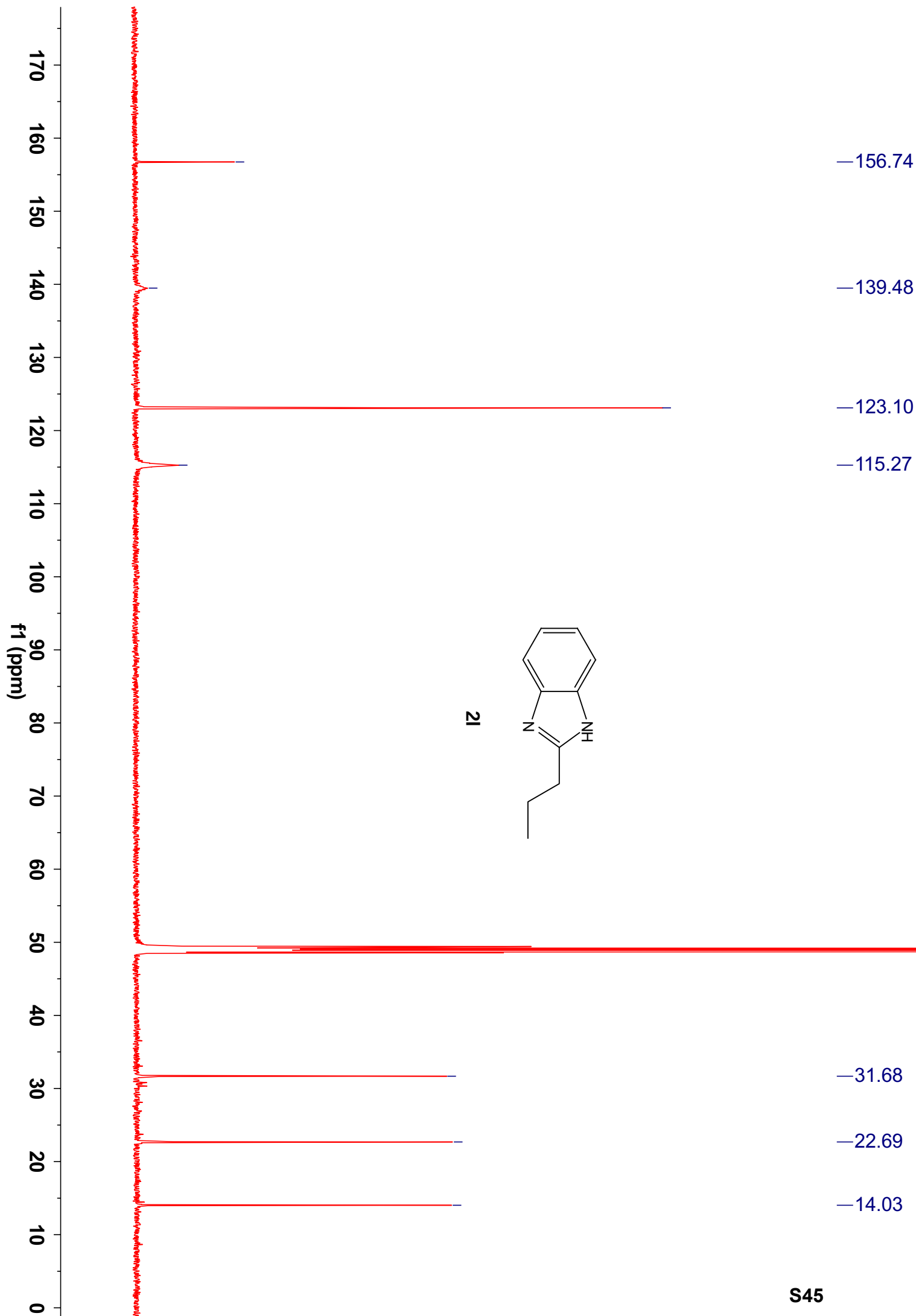
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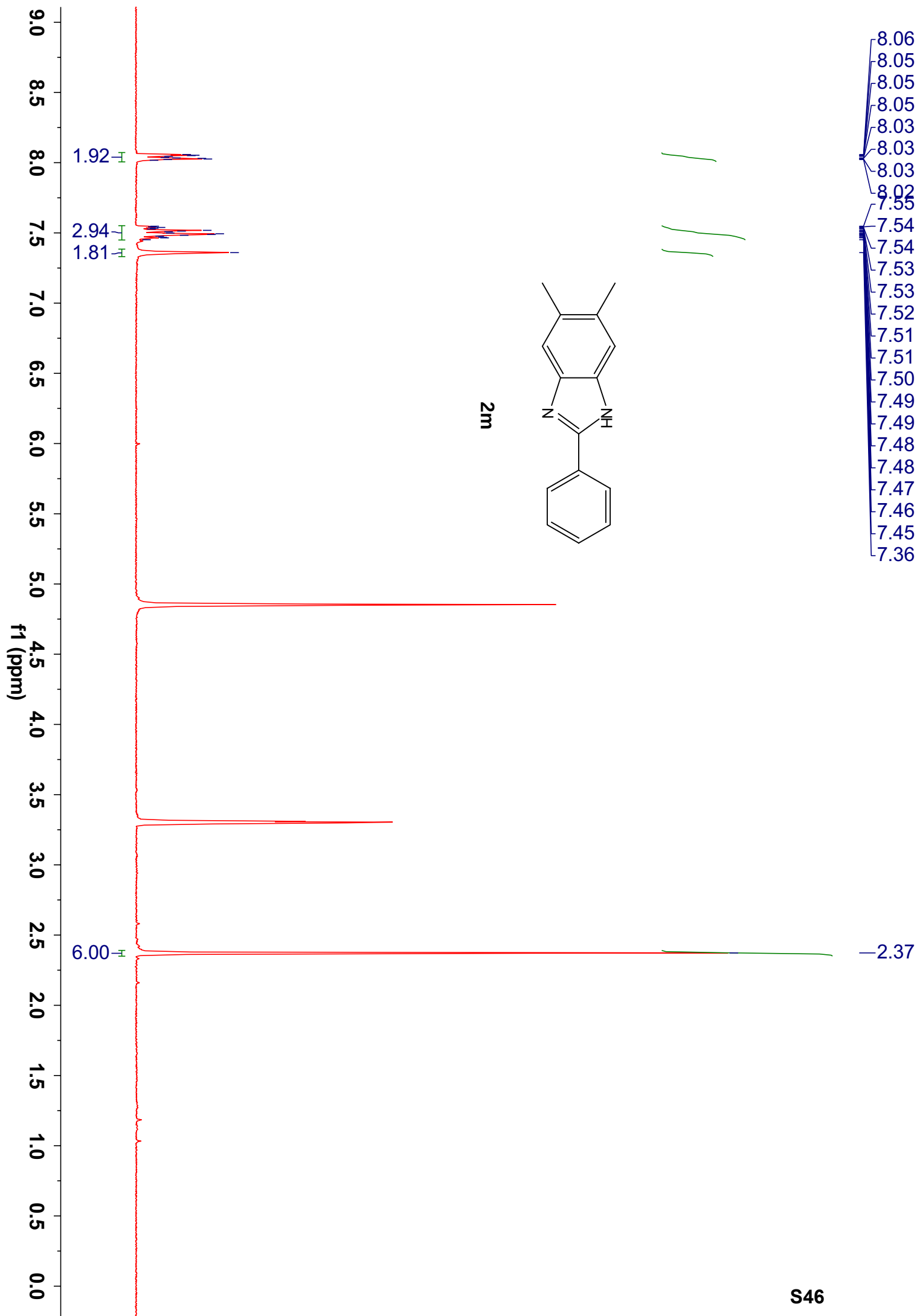
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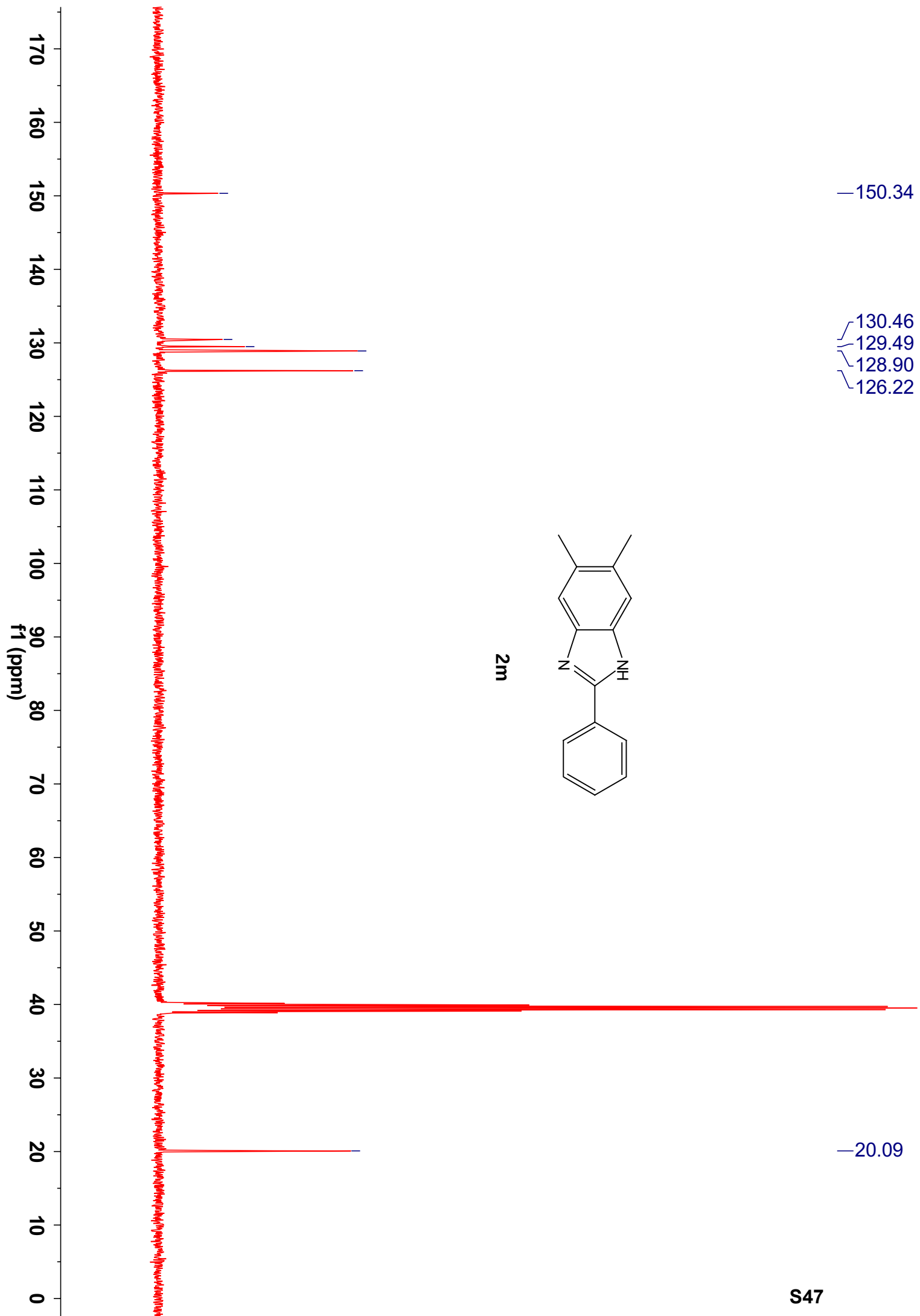


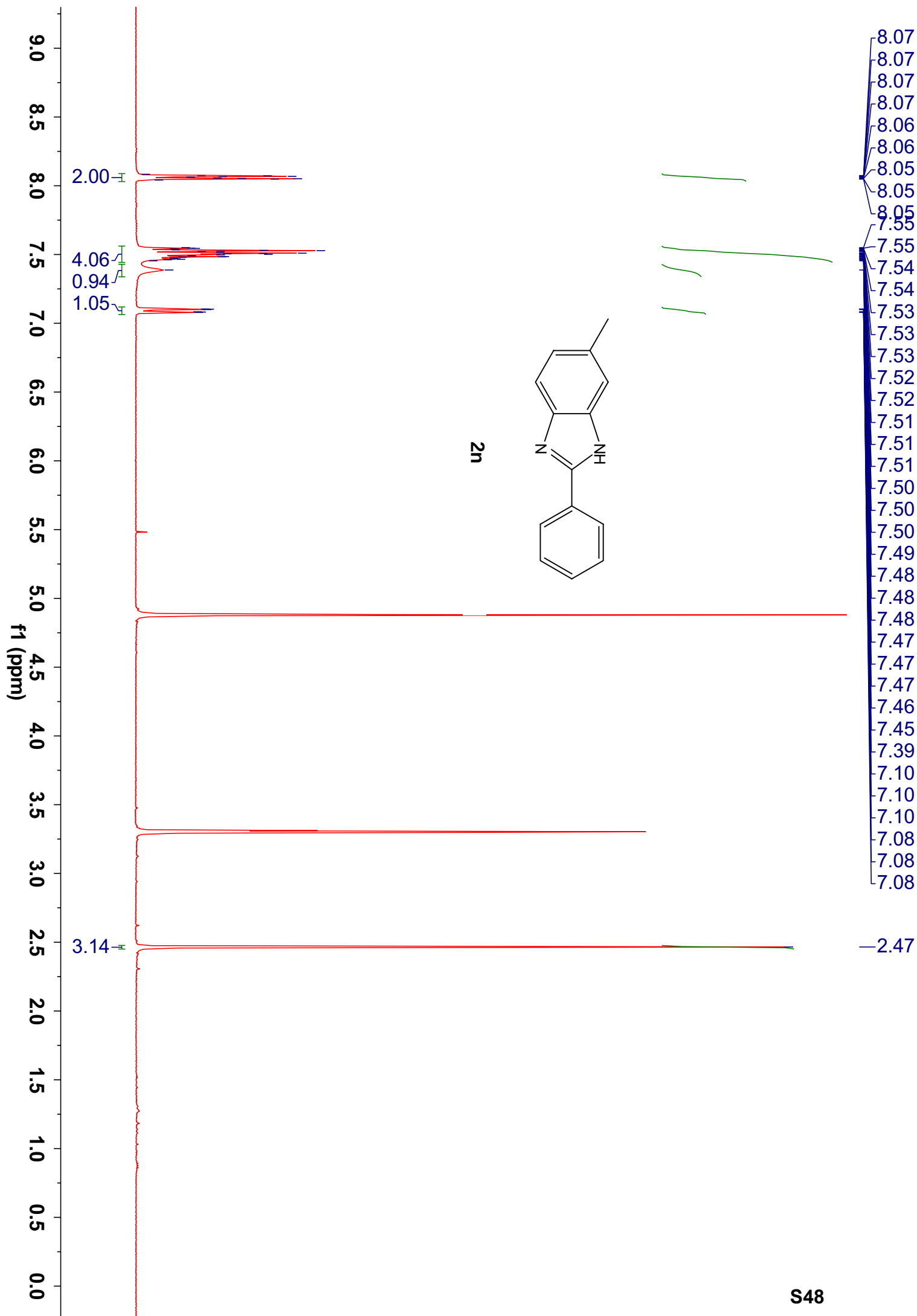
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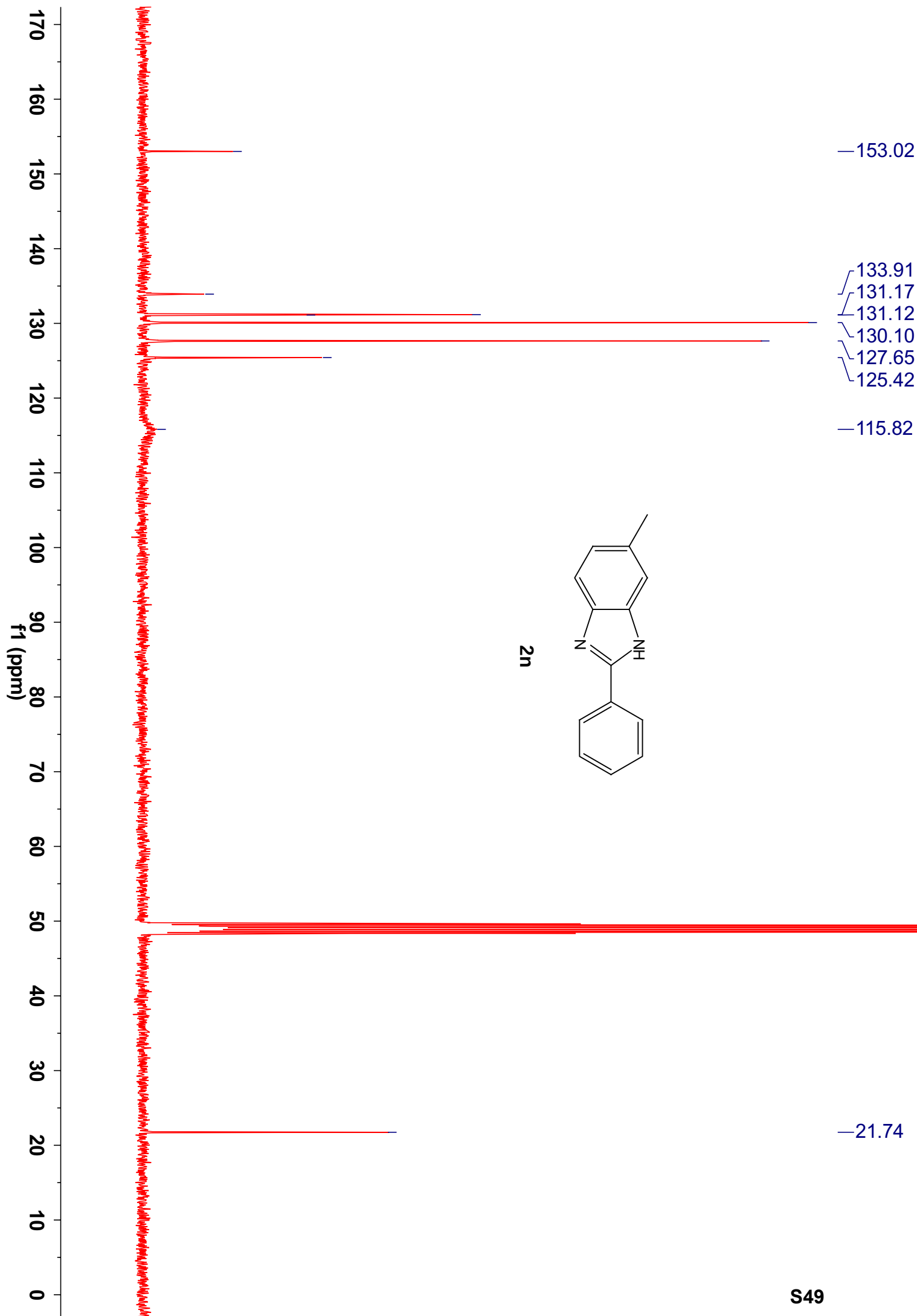


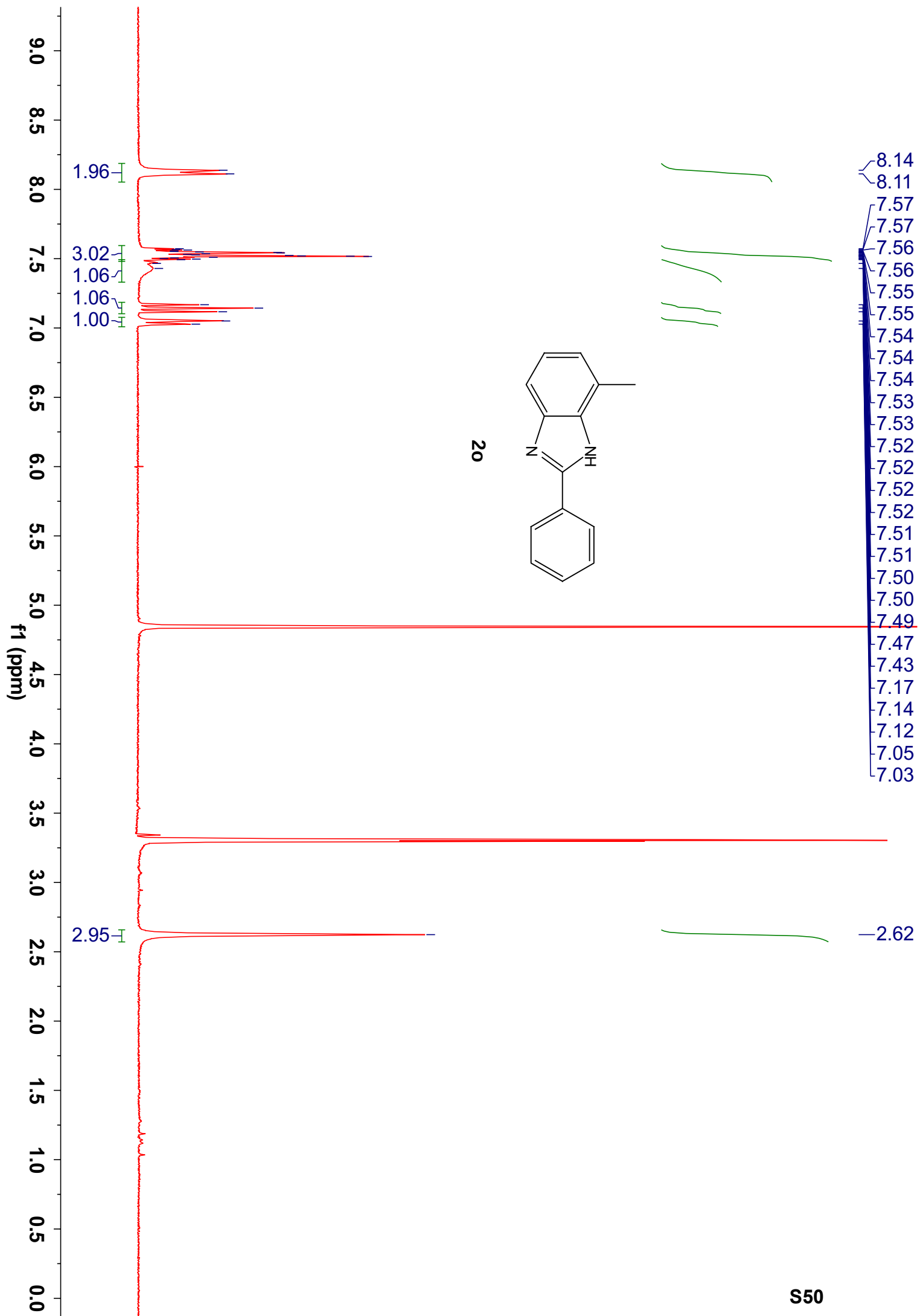


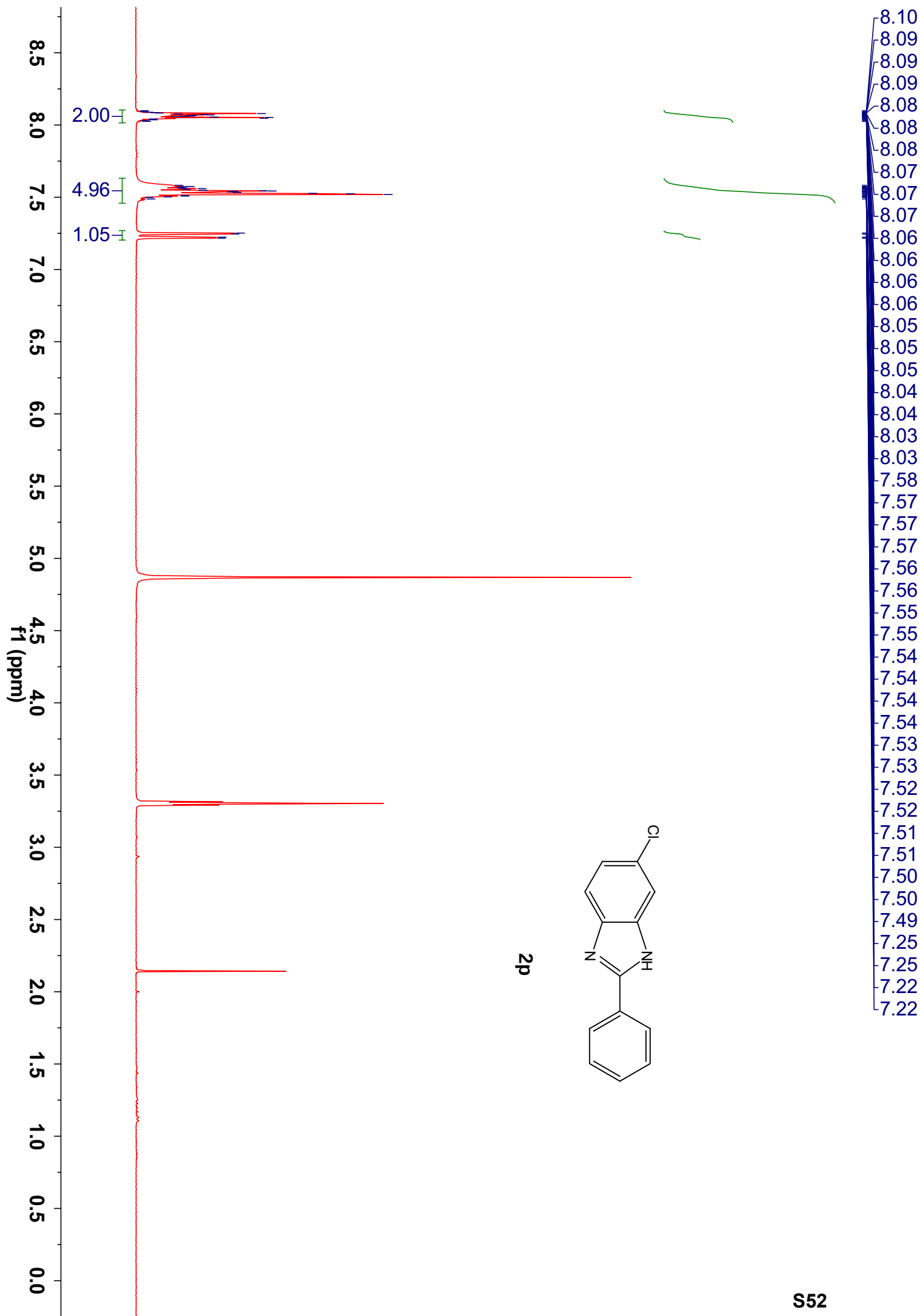


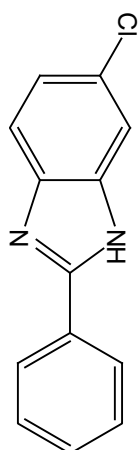




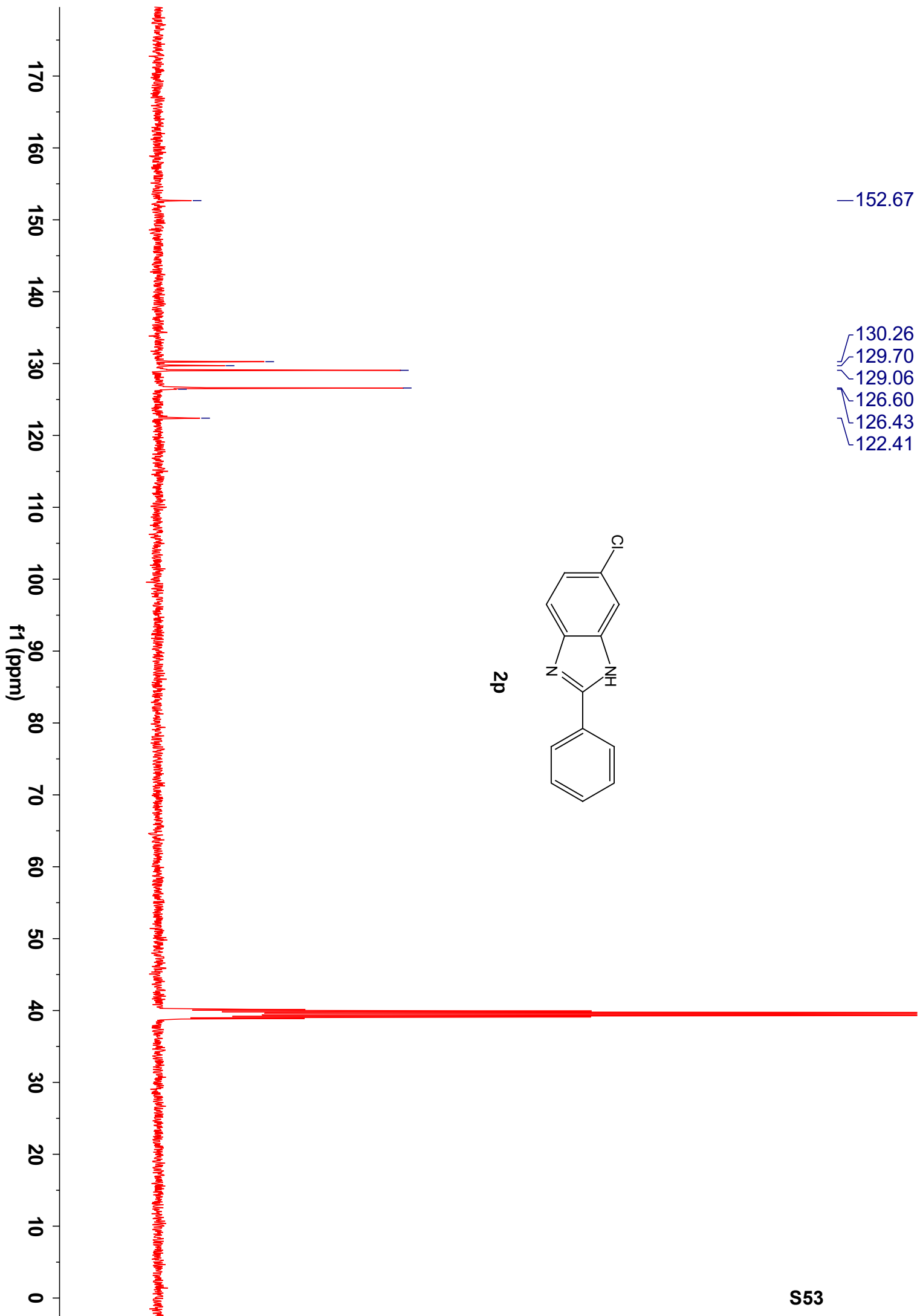


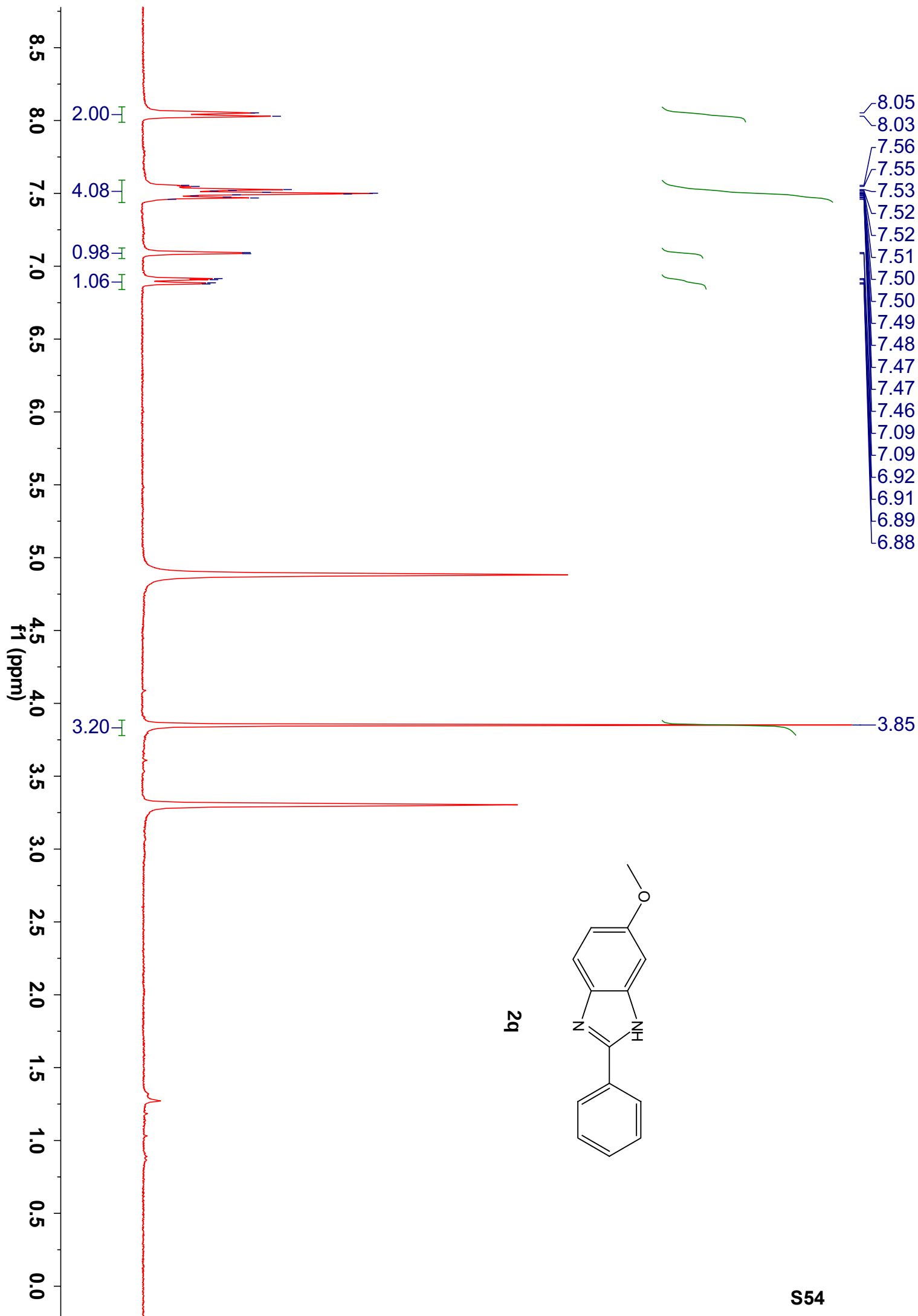


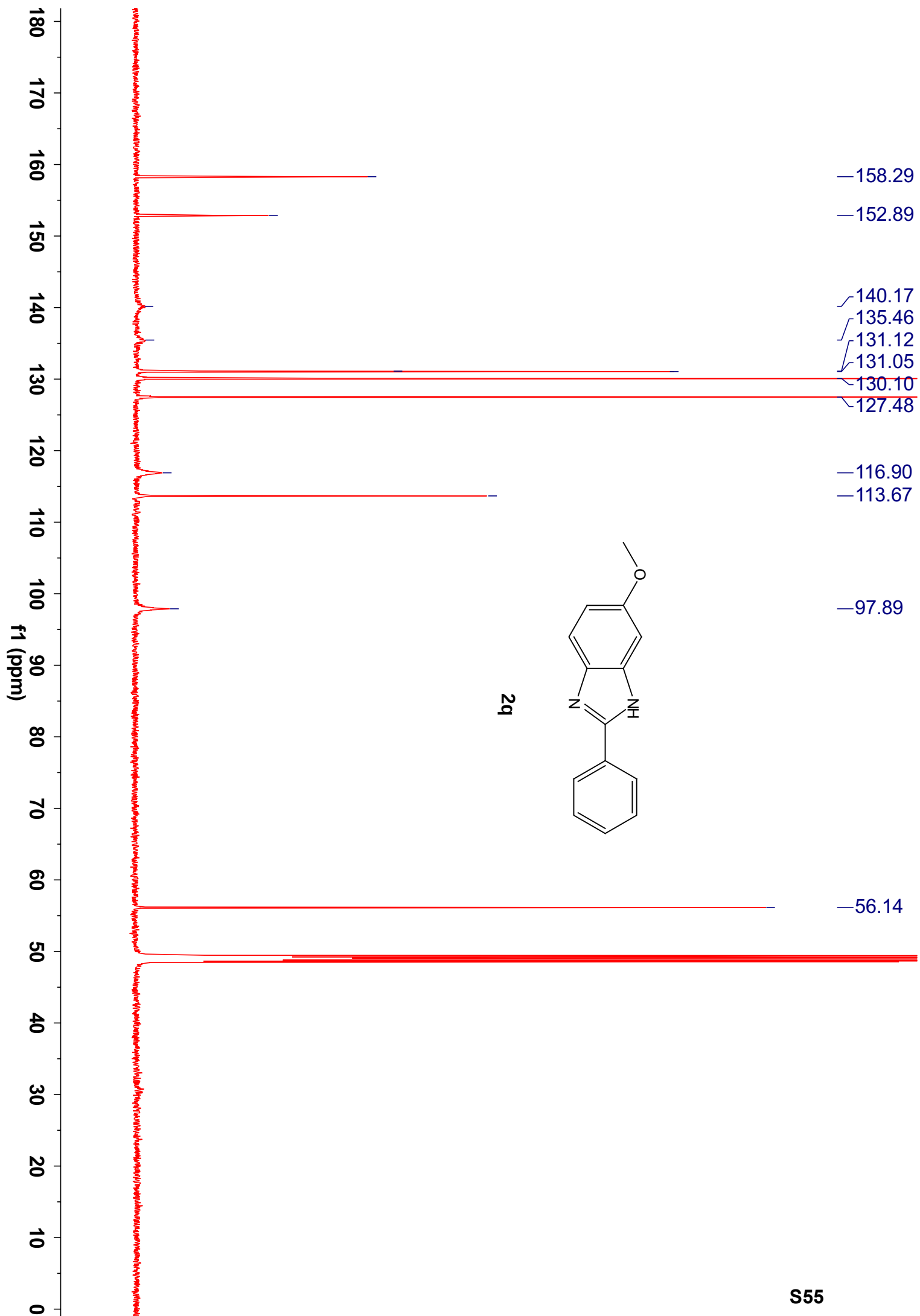


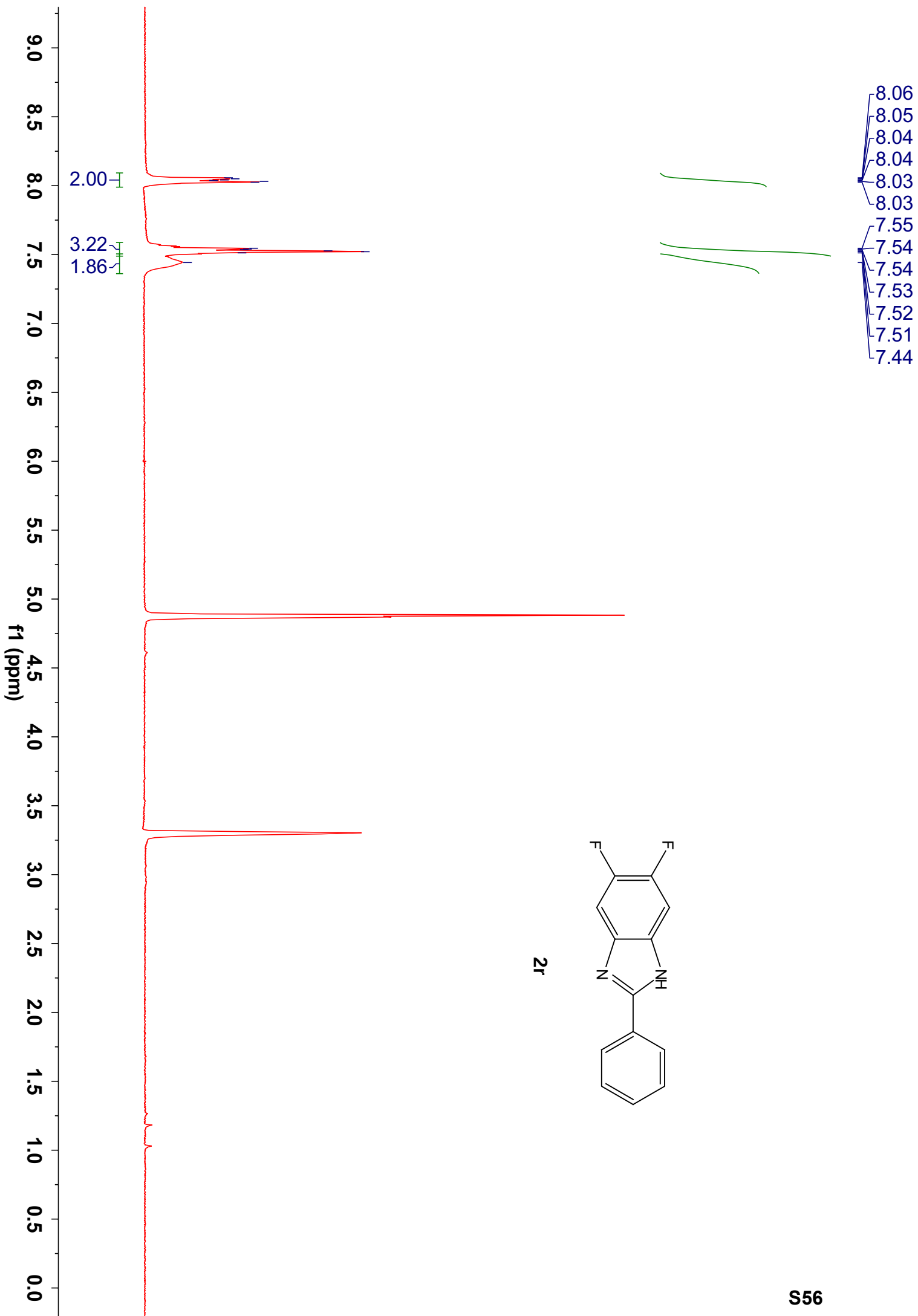
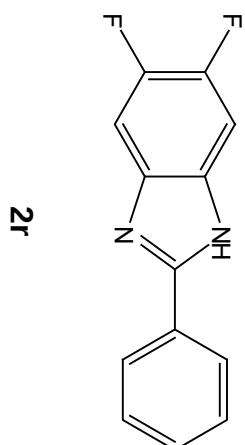


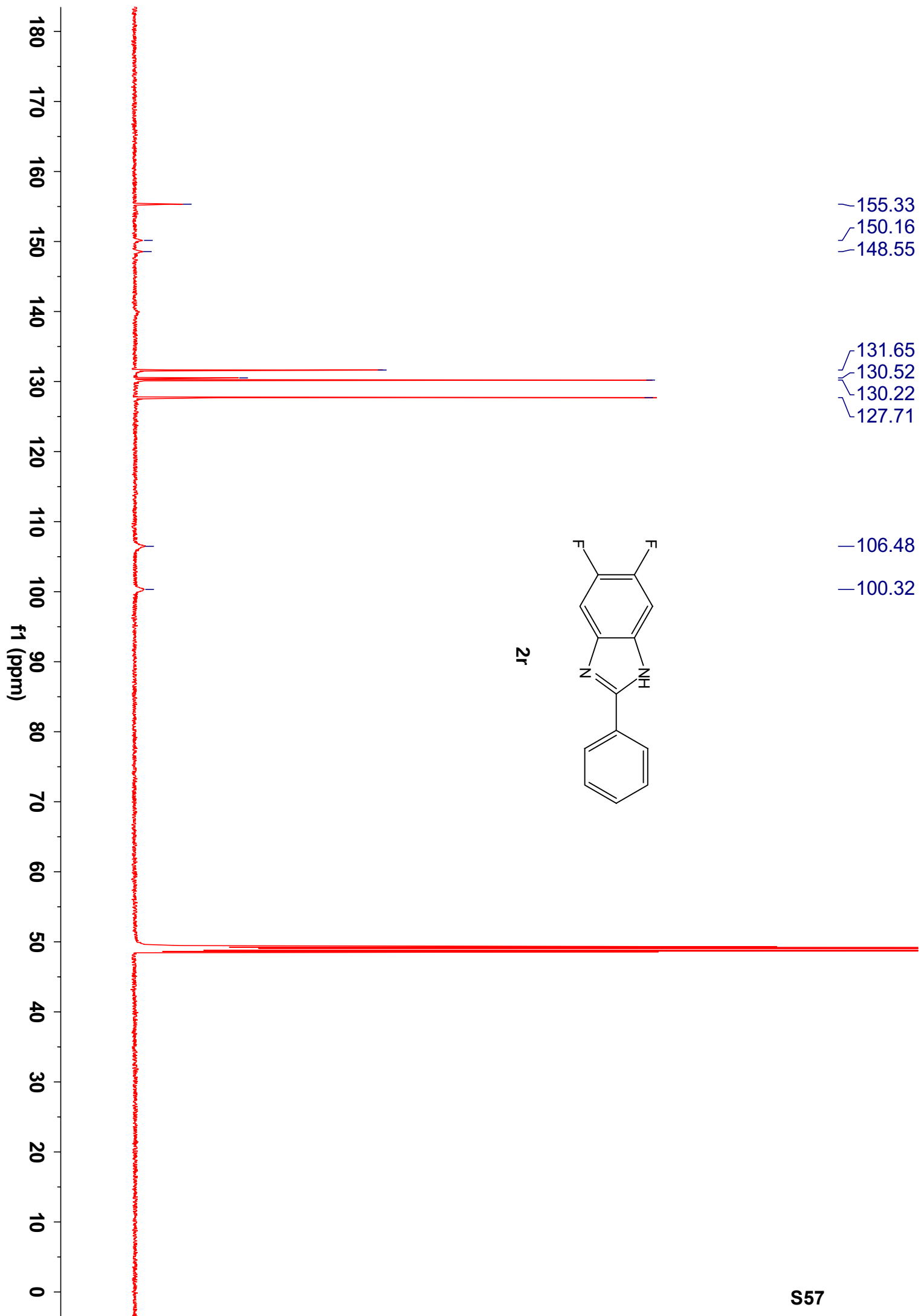
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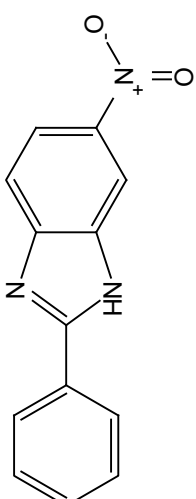




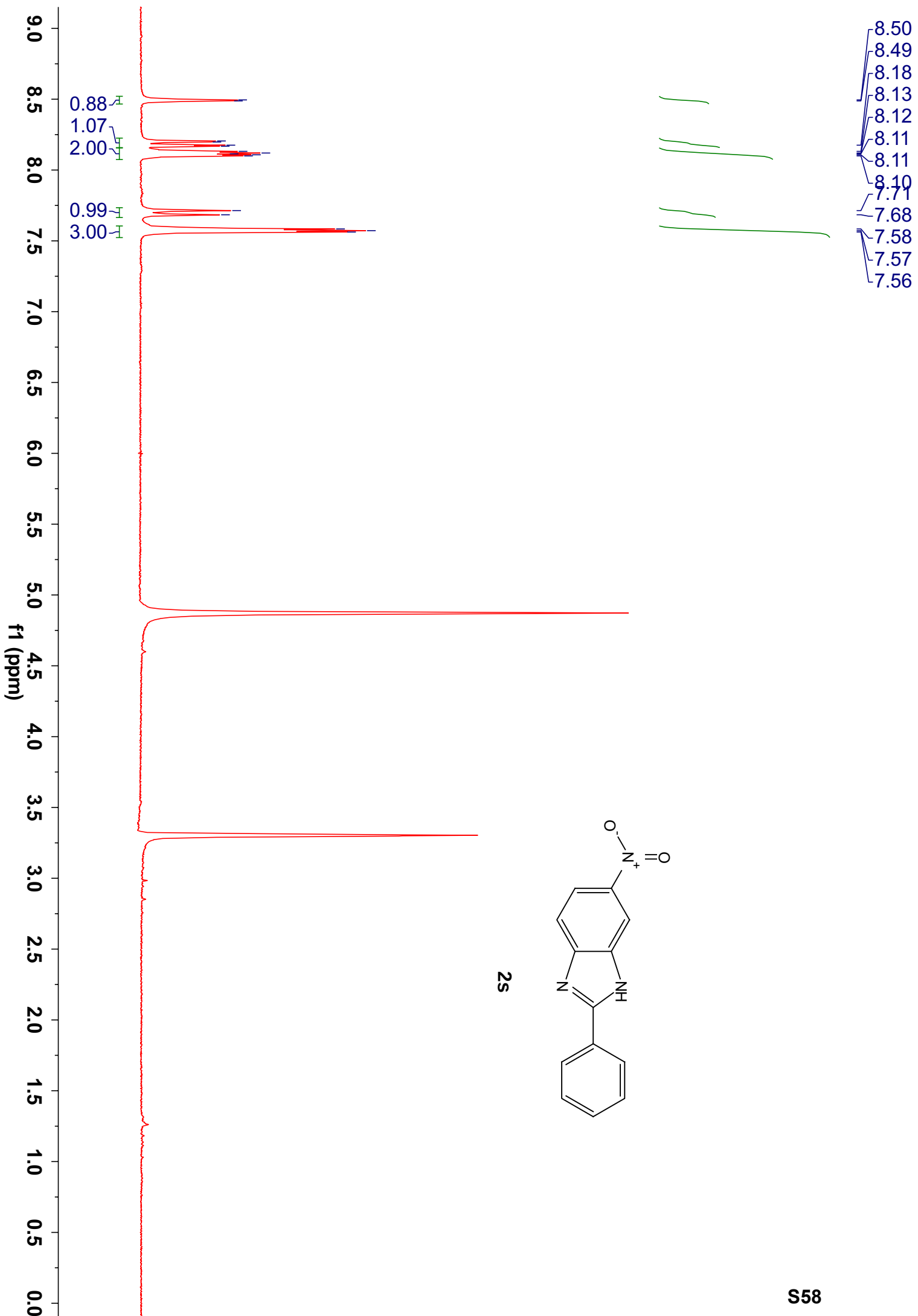


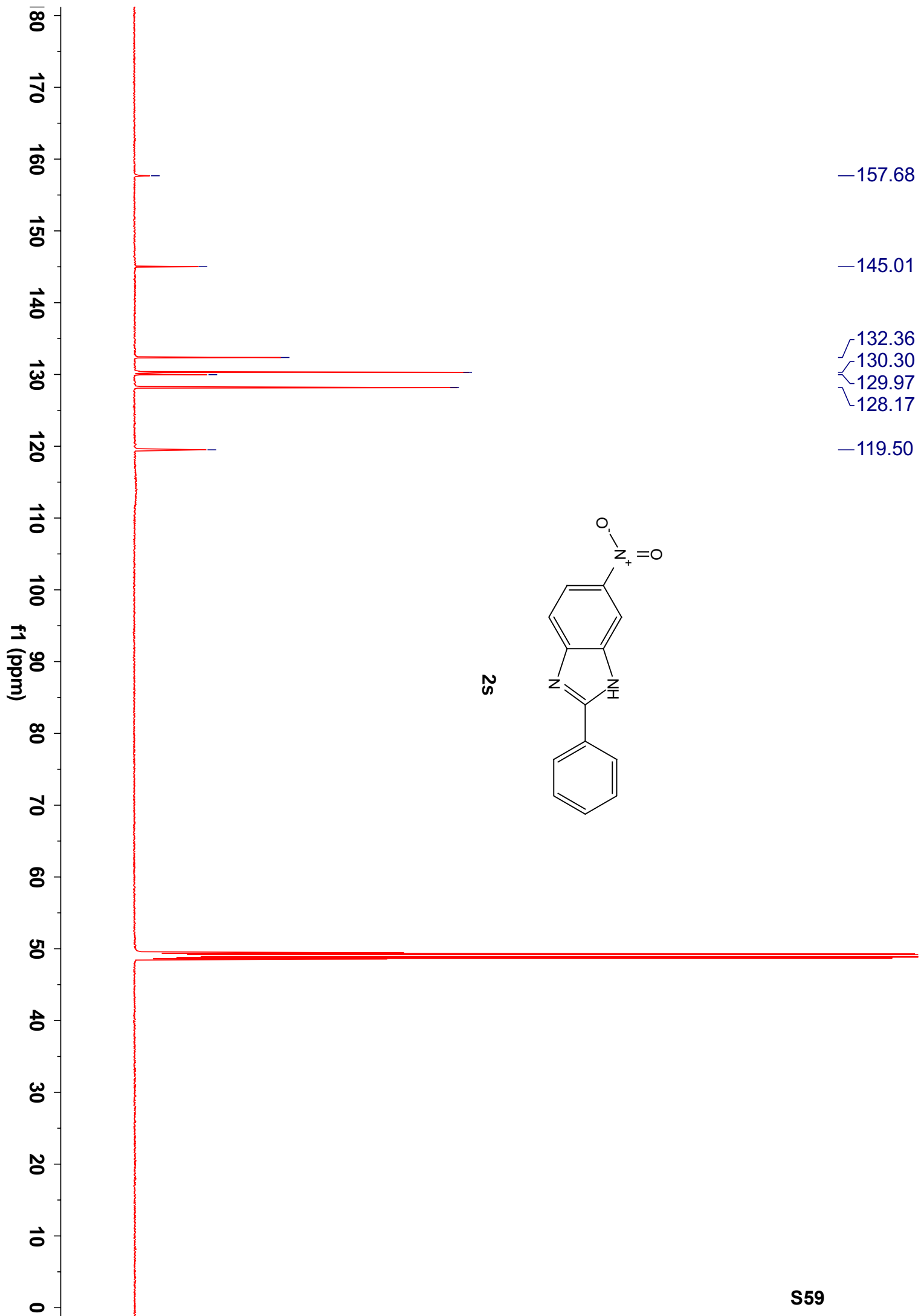


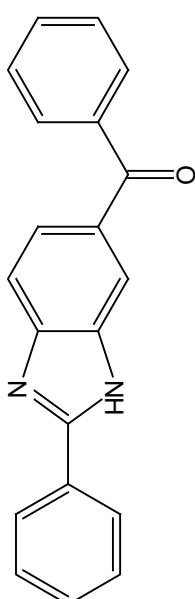
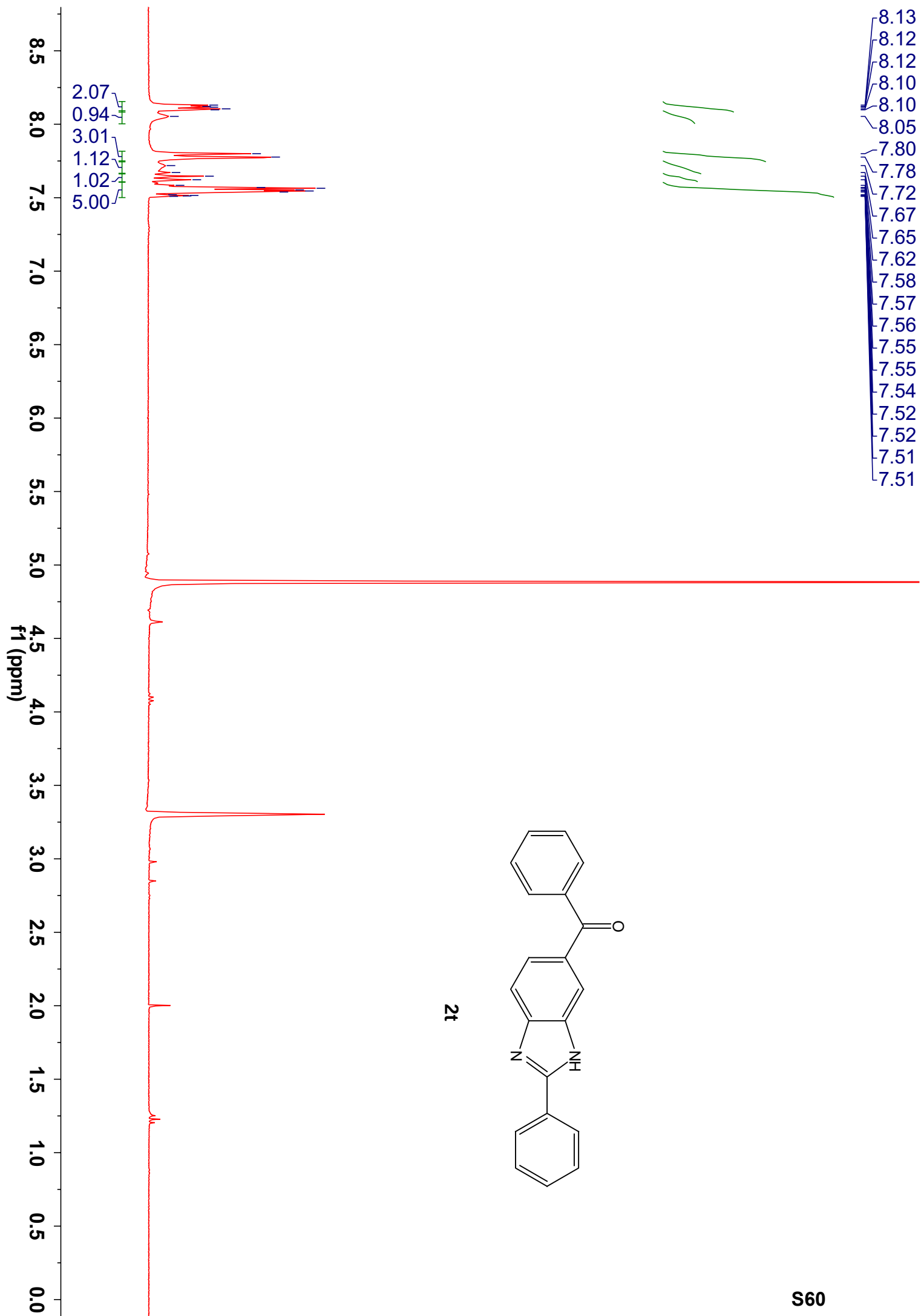


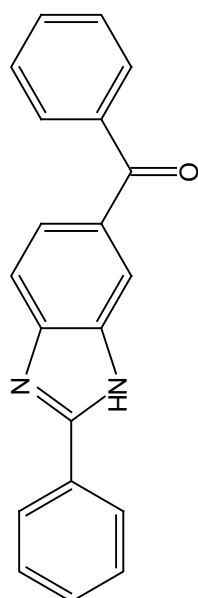


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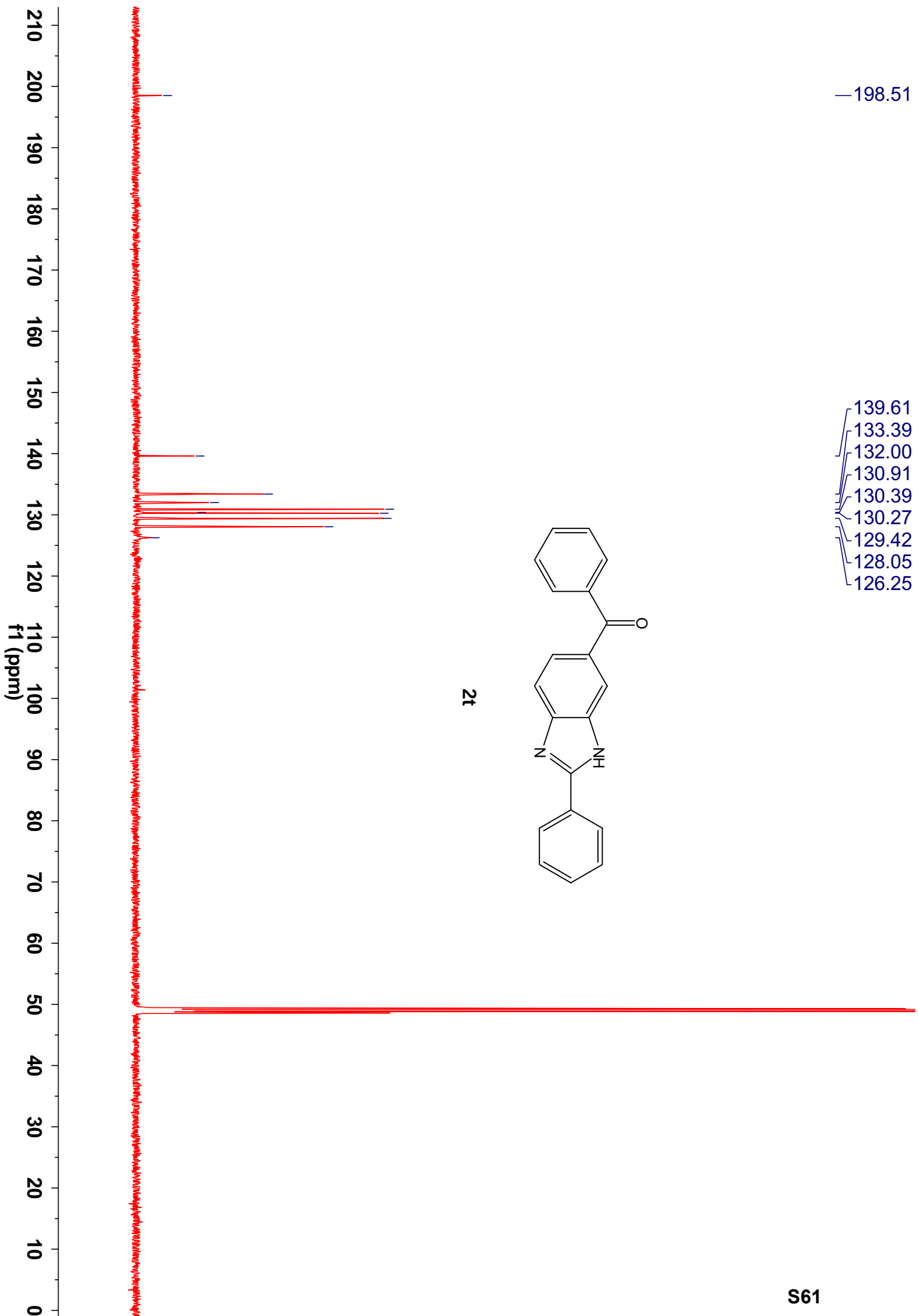


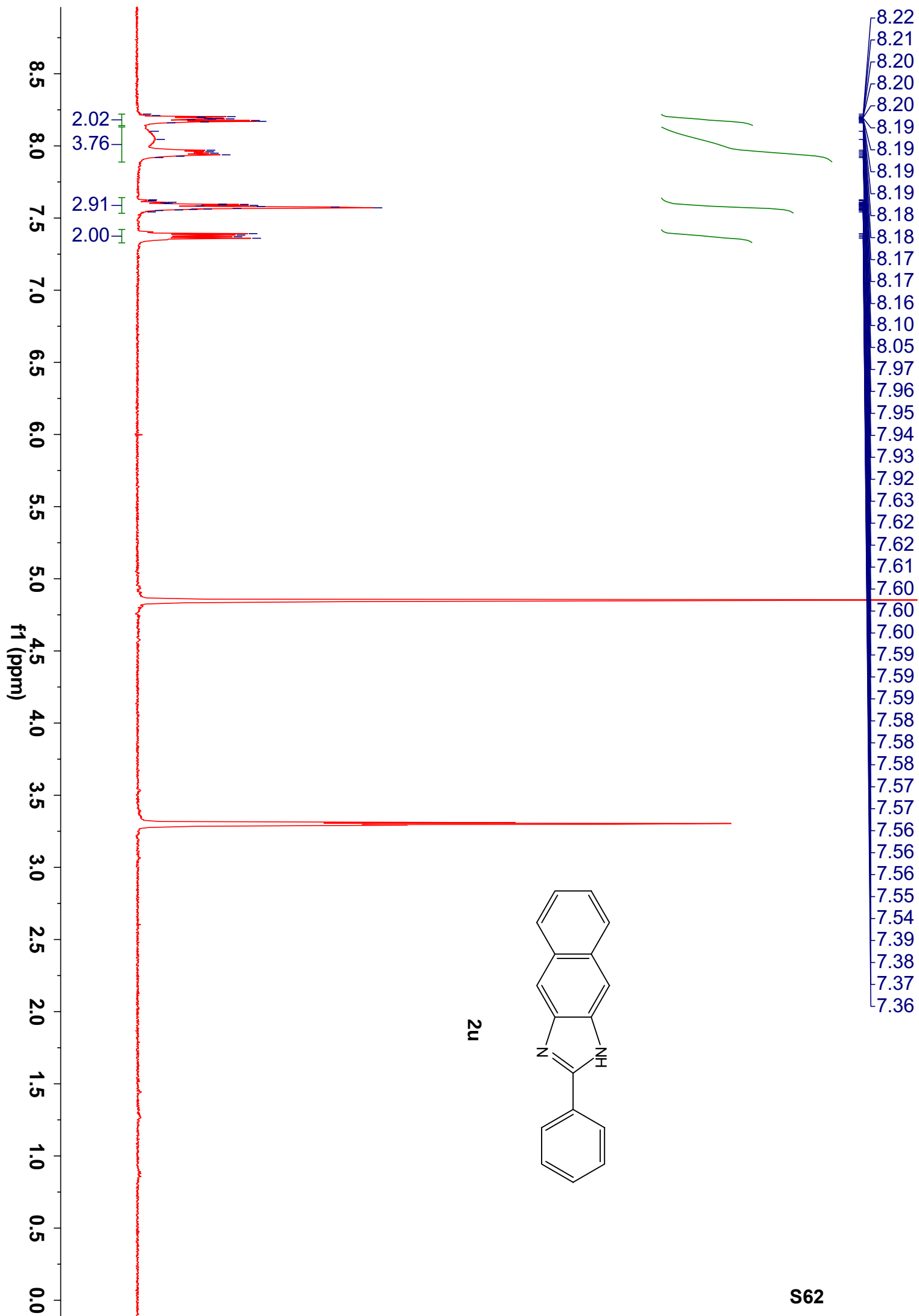


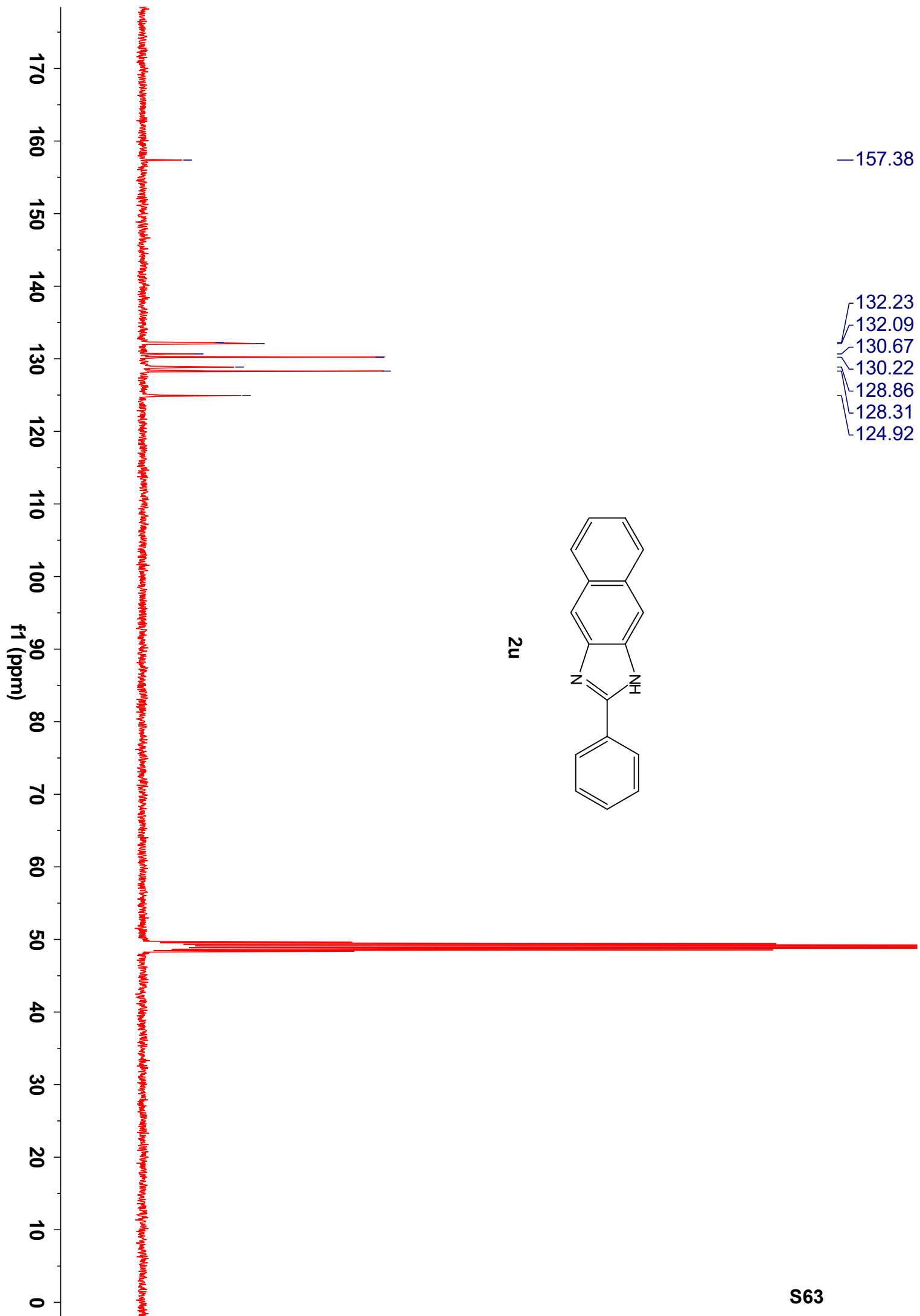
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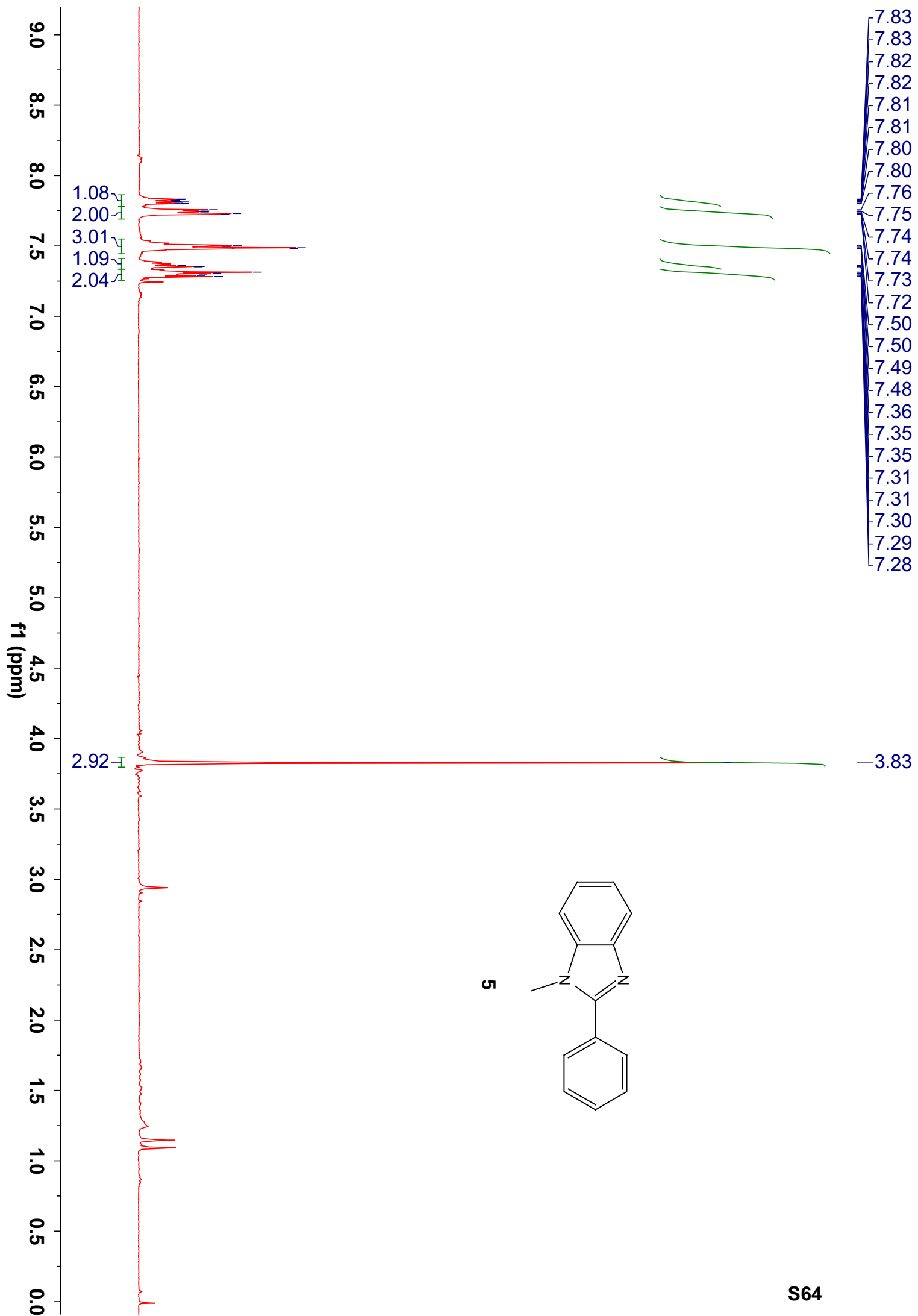


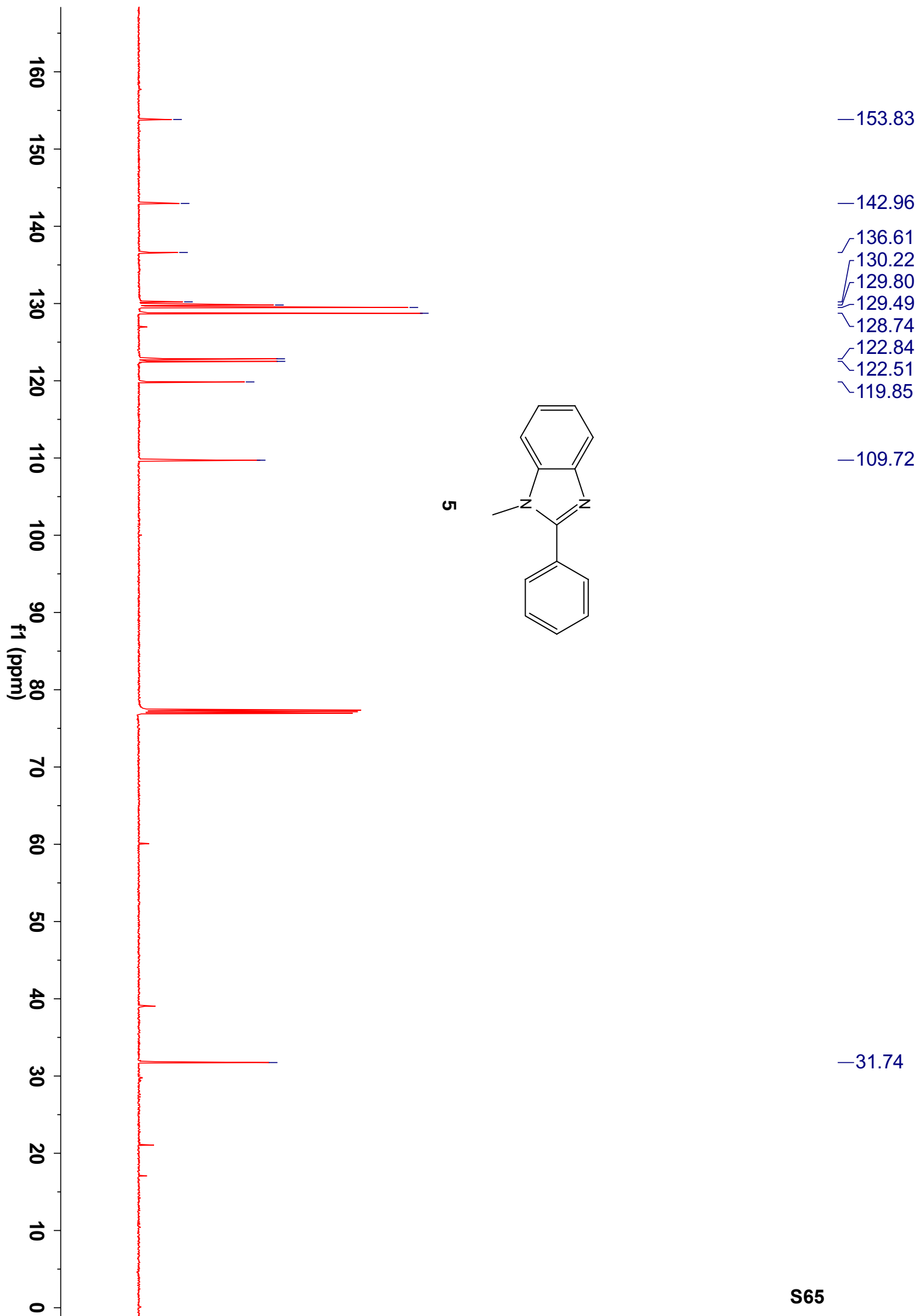
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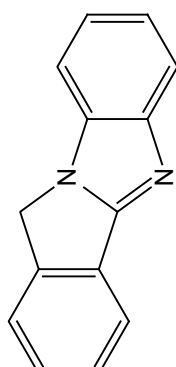




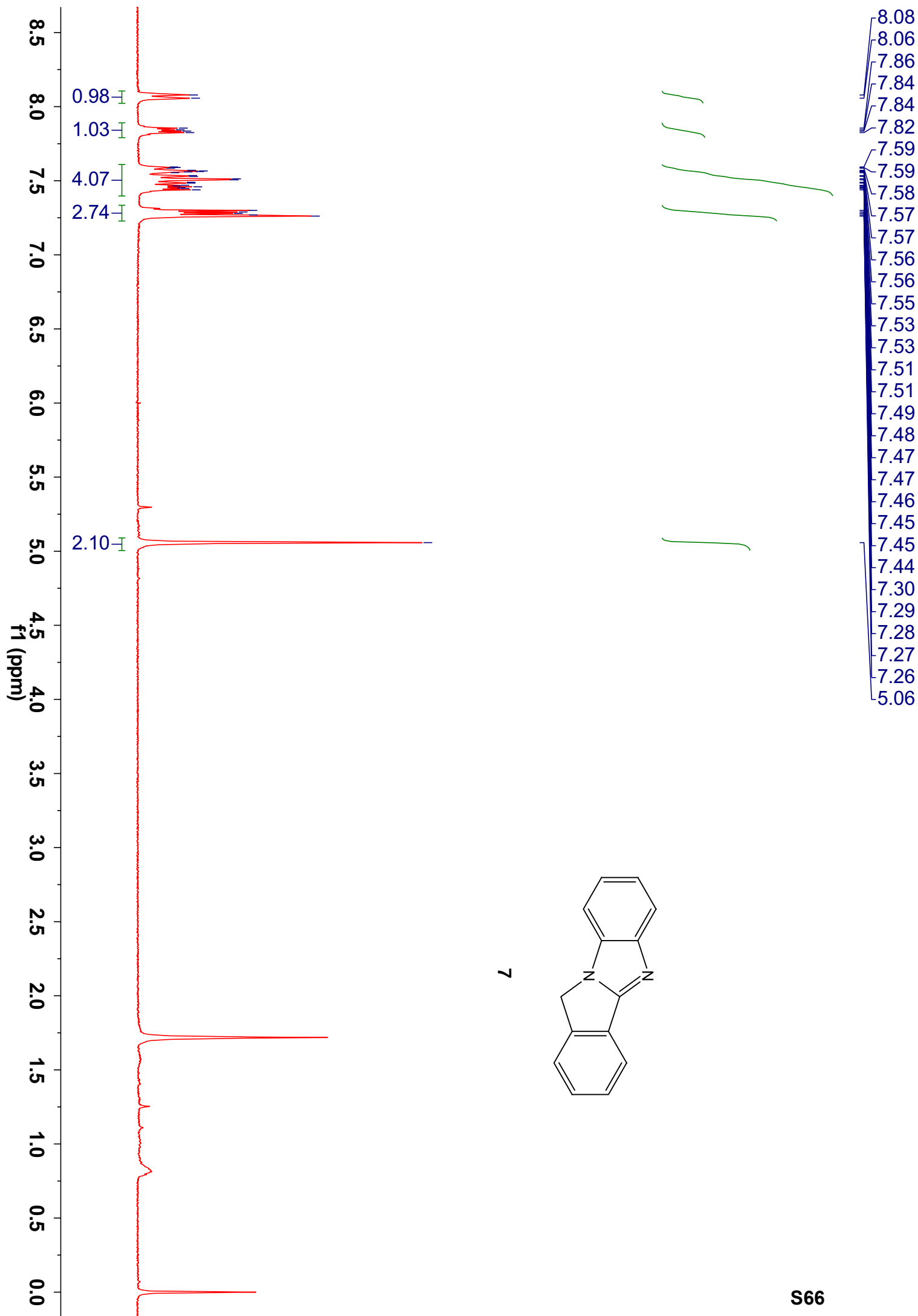


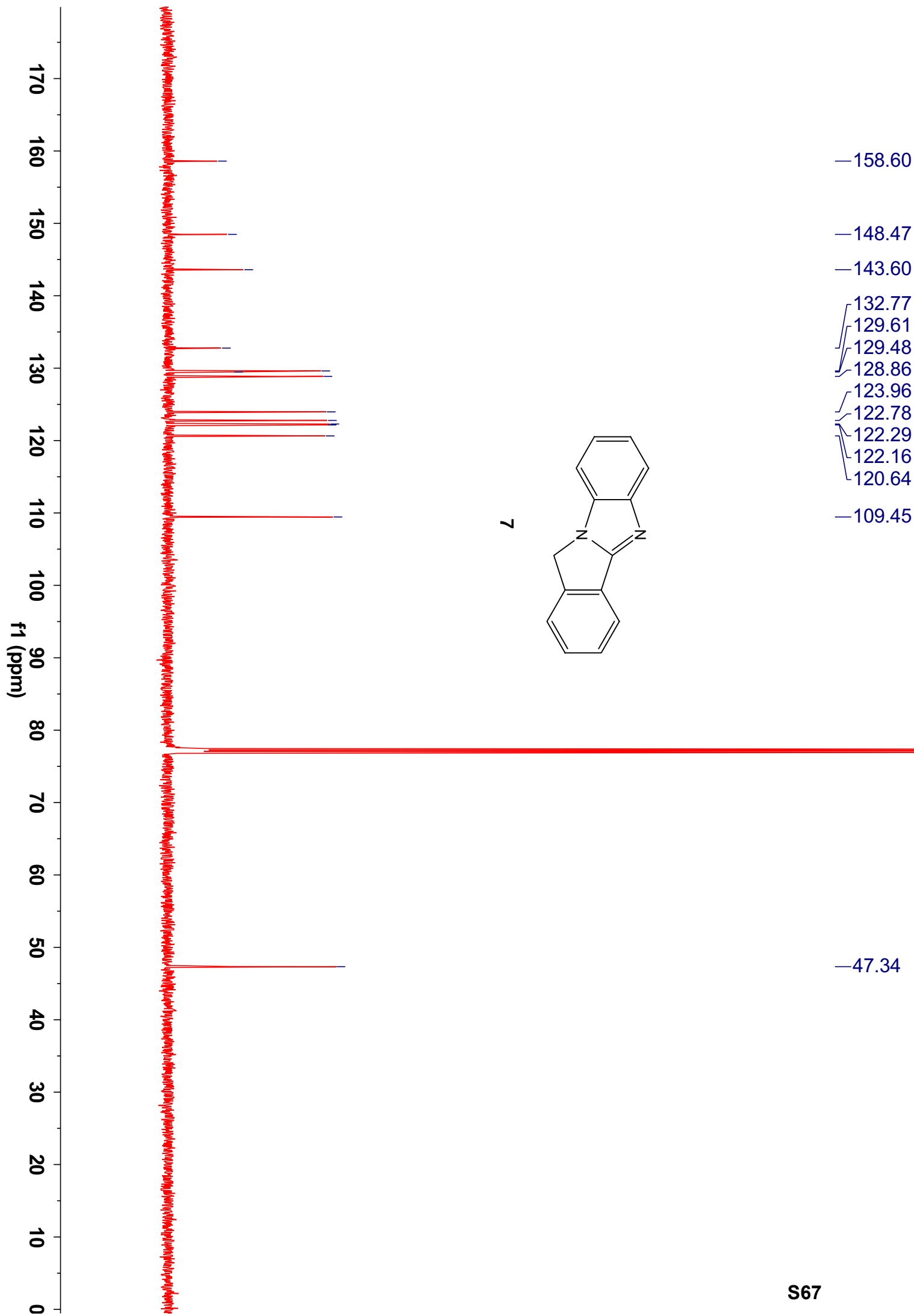


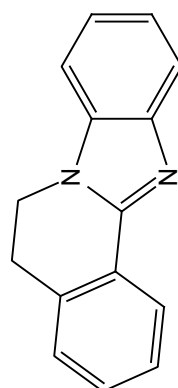




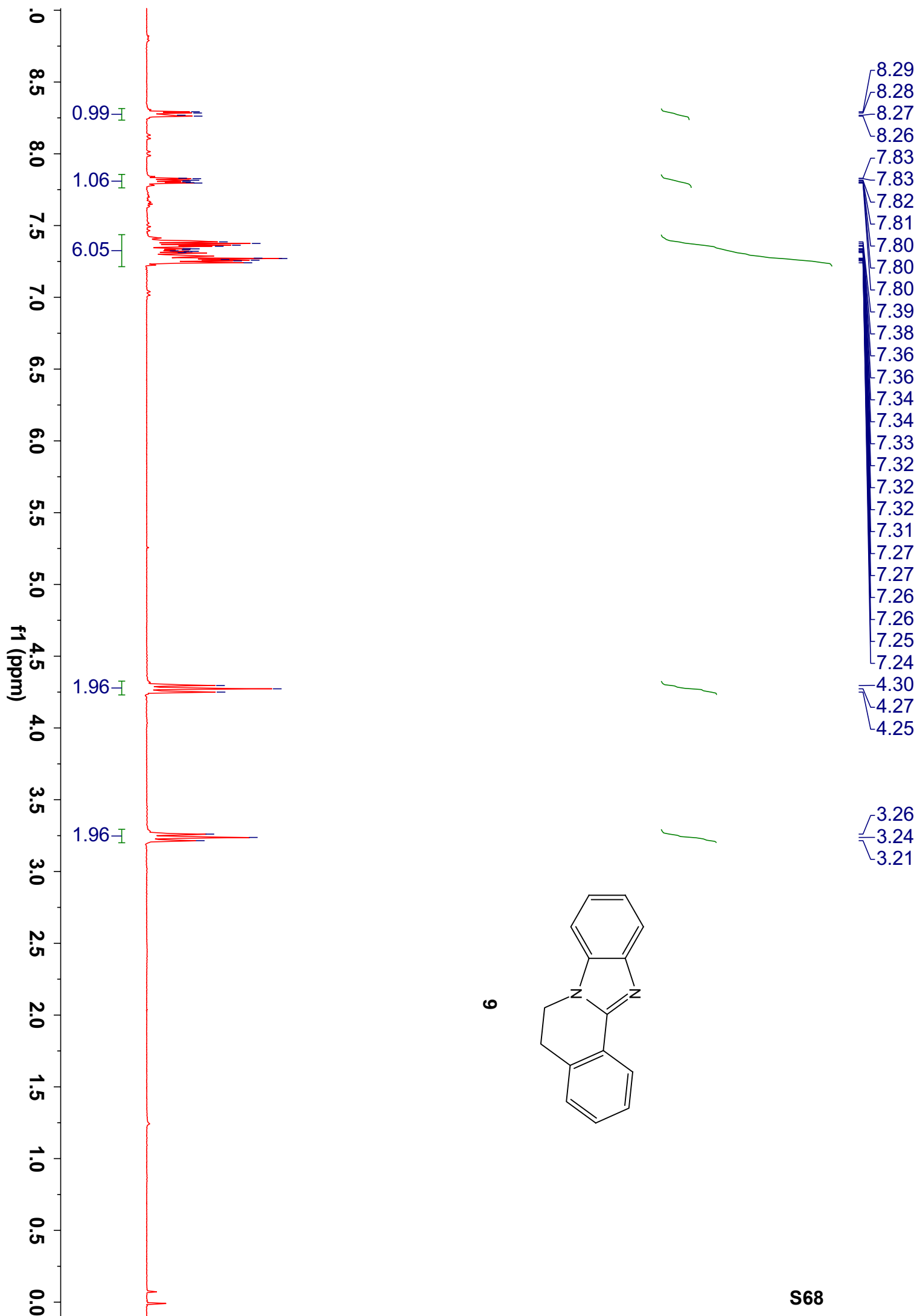
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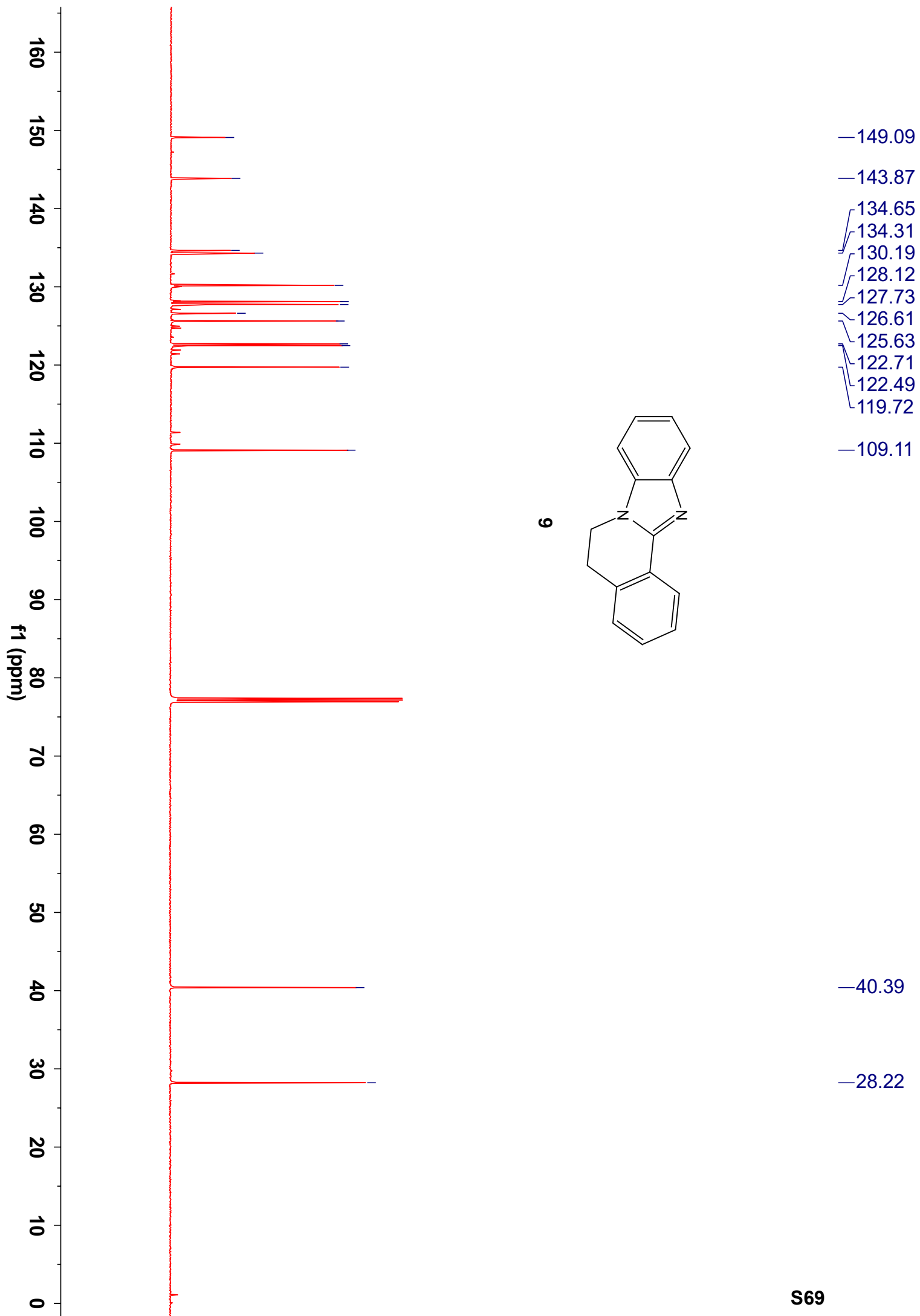
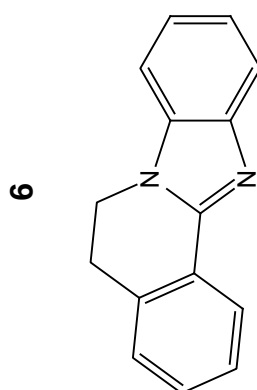


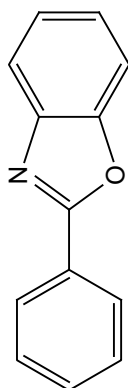




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**11**