

Supplementary Information for:

Regioselective β -Pyrrolic Electrophilic Substitution of Hydrodipyrin–Dialkylboron Complexes Facilitates Access to Synthetic Models for Chlorophyll *f*

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Table of Contents

Topic	Page
1. NMR Characterization of Dibutylboron Formyltetrahydrodipyrins	S1
2. X-ray Data	S2-S3
3. Decomplexation Results	S4
4. Spectral Data	S4-S43

1. NMR Characterization of Dibutylboron Formyltetrahydrodipyrins.

The resonances observed for protons in dibutylboron formyltetrahydrodipyrins **1-F⁷BBu₂**, **1-F⁸BBu₂** and **1-F⁹BBu₂** were assigned (Table S1). In **1-F⁹BBu₂** the formyl proton exhibited two cross peaks with a butyl group and H⁸. In **1-F⁸BBu₂** the formyl proton has two NOE with H⁹ and H⁷. And in **1-F⁷BBu₂** the formyl proton has two cross peaks with H⁸ and a methine proton. The observed NOE of the pyrrolic protons in formyltetrahydrodipyrin–dibutylboron complexes also strongly supported the assigned structures (Figure S1).

Table S1. ¹H NMR chemical shifts of dibutylboron formyltetrahydrodipyrins.

Compound	H-formyl	H ⁷	H ⁸	H ⁹
1-F⁷BBu₂	9.77	—	6.55	6.62
1-F⁸BBu₂	9.68	6.37	—	7.29
1-F⁹BBu₂	9.81	6.02	7.14	—

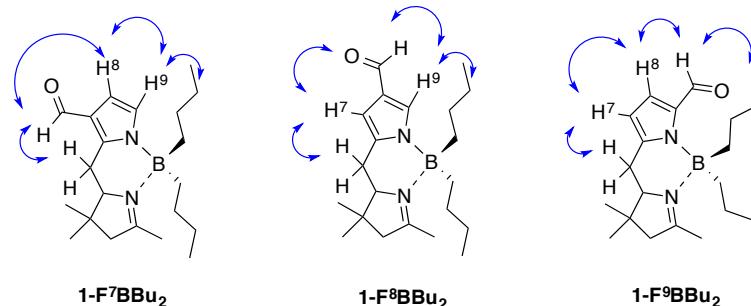


Figure S1. NOEs observed with tetrahydrodipyrin–dialkylboron complexes.

The position of the formyl group in **1-F⁷BBu₂** was established by ¹H NMR spectroscopy with observation of the following NOE signals: (1) the NOE signal between the formyl proton and one of the methylene group protons (note the chemical shift change of one of the methylene group protons is pronounced upon formylation, from 2.82–2.85 ppm to 3.57–3.63 ppm); (2) the NOE signal between the formyl proton and the β -pyrrolic proton; (3) the NOE signal between the α -pyrrolic proton and the butyl group; and (4) the NOE signal between the α - and β -pyrrolic protons.

2. X-ray Data.

Table S2. Summary of Crystal Data for **1-BBu₂**, **1-Br⁸BBu₂**, **2-BBu₂**, and **1-Br⁸F⁷BBu₂**.

	1-BBu₂	1-Br⁸BBu₂	2-BBu₂	1-Br⁸F⁷BBu₂
CCDC registry	974409	974410	974412	974411
Formula	C ₂₀ H ₃₅ BN ₂	C ₂₀ H ₃₄ BBrN ₂	C ₂₇ H ₃₉ BN ₂	C ₂₁ H ₃₄ BBrN ₂ O
Formula Weight (<i>g/mol</i>)	314.32	393.21	402.41	421.22
Crystal Dimensions (mm)	0.32 × 0.20 × 0.09	0.45 × 0.30 × 0.20	0.36 × 0.26 × 0.08	0.50 × 0.37 × 0.09
Crystal System	monoclinic	monoclinic	monoclinic	orthorhombic
Space Group	<i>P</i> 2 ₁ / <i>n</i>	<i>P</i> 2 ₁ / <i>c</i>	<i>P</i> 2 ₁ / <i>c</i>	<i>Pna</i> 2 ₁
Temperature, K	110	110	173	110
<i>a</i> , Å	15.8194(3)	8.8715(3)	9.077(2)	17.5208(3)
<i>b</i> , Å	7.93900(10)	26.3874(12)	16.466(6)	16.5255(3)
<i>c</i> , Å	16.6596(3)	9.8079(4)	18.163(6)	15.1765(3)
α, deg	90	90	90	90
β, deg	110.7227(8)	115.588(2)	109.714(8)	90
γ, deg	90	90	90	90
V, Å ³	1956.92(6)	2070.80(15)	2555.6(14)	4394.20(14)
Number of reflections to determine final unit cell	9052	503	8126	9969
Min and Max 2θ for cell determination (deg)	4.4, 44.38	4.85, 51.95	4.76, 42.28	5.26, 62.01
Z	4	4	4	8
F(000)	696.32	832	880	1776
ρ (g/cm)	1.067	1.261	1.046	1.273
λ, Å, (MoKα)	0.71073	0.71073	0.71073	0.71073
μ, (cm ⁻¹)	0.06	1.989	0.06	1.883
Max 2θ for data collection (deg)	53.32	56.56	52.82	54.64
Measured fraction of data	0.999	0.999	0.999	0.999
Number of reflections measured	110726	29325	54612	77149
Unique reflections measured	4114	5130	5249	9523
R _{merge}	0.031	0.0377	0.0461	0.0377
Number of parameters in least-squares	349	222	277	500
R ₁	0.052	0.0643	0.0696	0.0456
wR ₂	0.079	0.1295	0.191	0.1098
R ₁ (all data)	0.074	0.0771	0.1097	0.0497
wR ₂ (all data)	0.157	0.1343	0.222	0.1119

Compound **1-Br⁸F⁷BBu₂** crystallizes in the chiral space group. The absolute structure parameter for **1-Br⁸F⁷BBu₂** was refined using TWIN/BASF commands to accommodate for a small amount of racemic twinning. Electron density peaks in the difference map of **1-Br⁸F⁷BBu₂** revealed disorder in the position of the bromine atom (Br1, Br1A; Br2, Br2A) on both of the molecules in the asymmetric unit (at the respective positions C8, C9 and C29, C30). The two enantiomers of **1-Br⁸F⁷BBu₂** are displayed in Figure S2. In addition, the two ball-and-stick models show the composite electron density for the presence of **1-Br⁸F⁷BBu₂** and **1-Br⁹F⁷BBu₂** in 97:3 ratio.

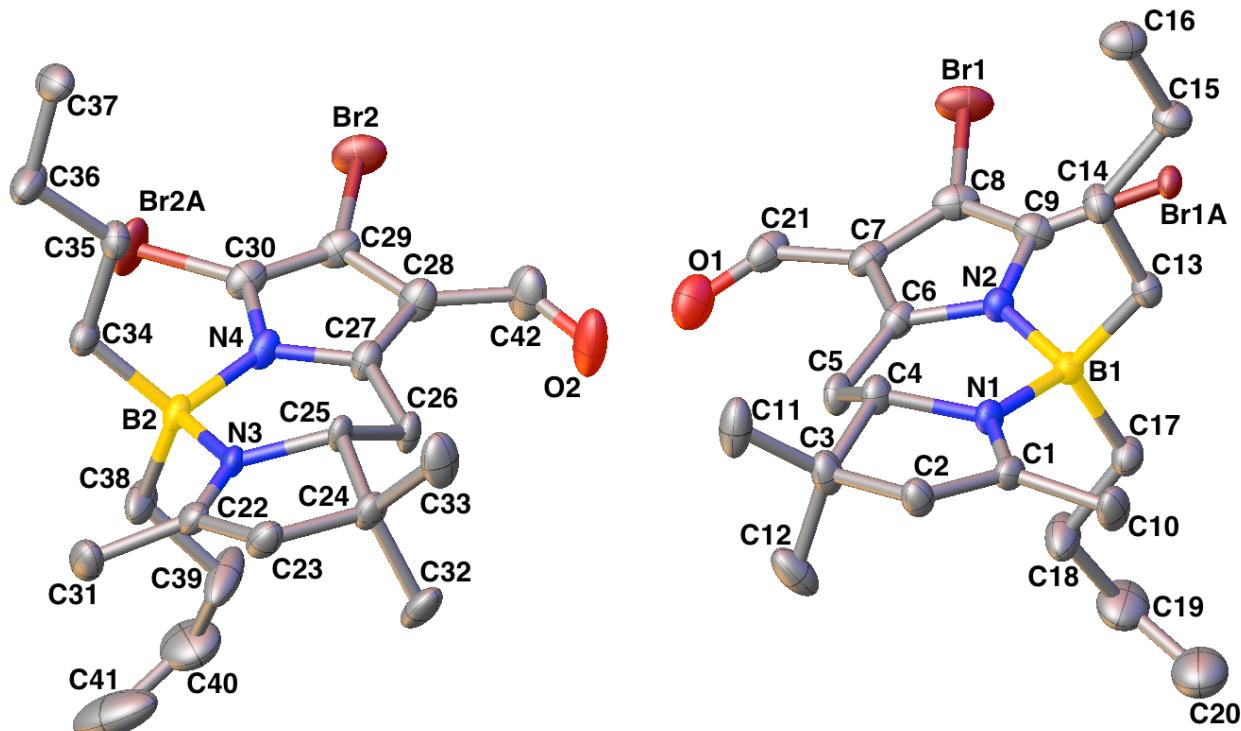


Figure S2. Enantiomers of **1-Br⁸F⁷BBu₂** also showing the disorder of the bromo position.

3. Decomplexation Results

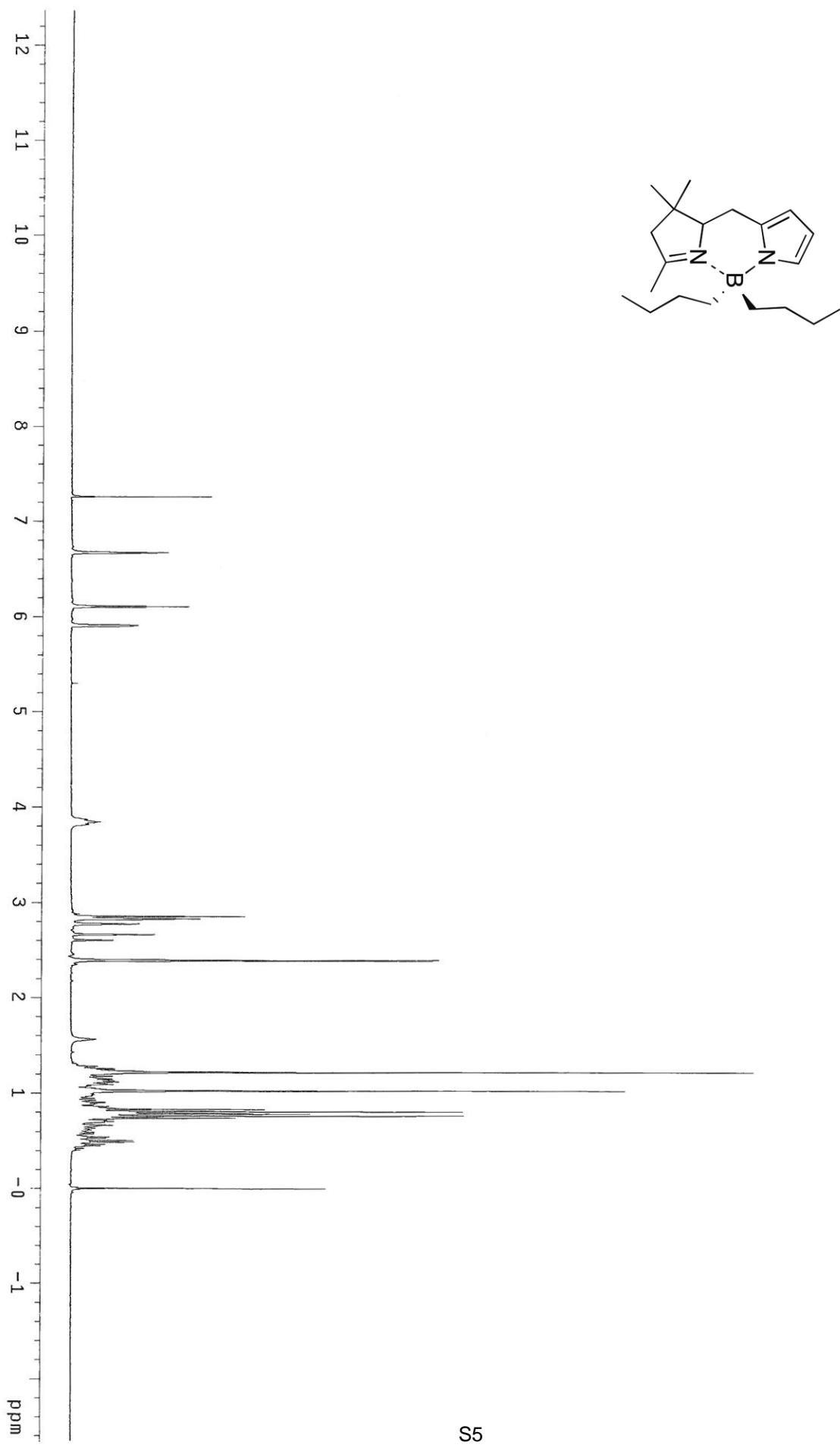
Table S3. Conditions for decomplexation of **1-BR₂**.

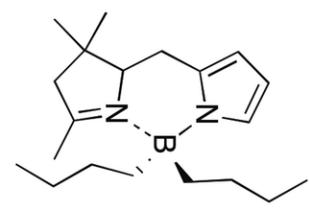
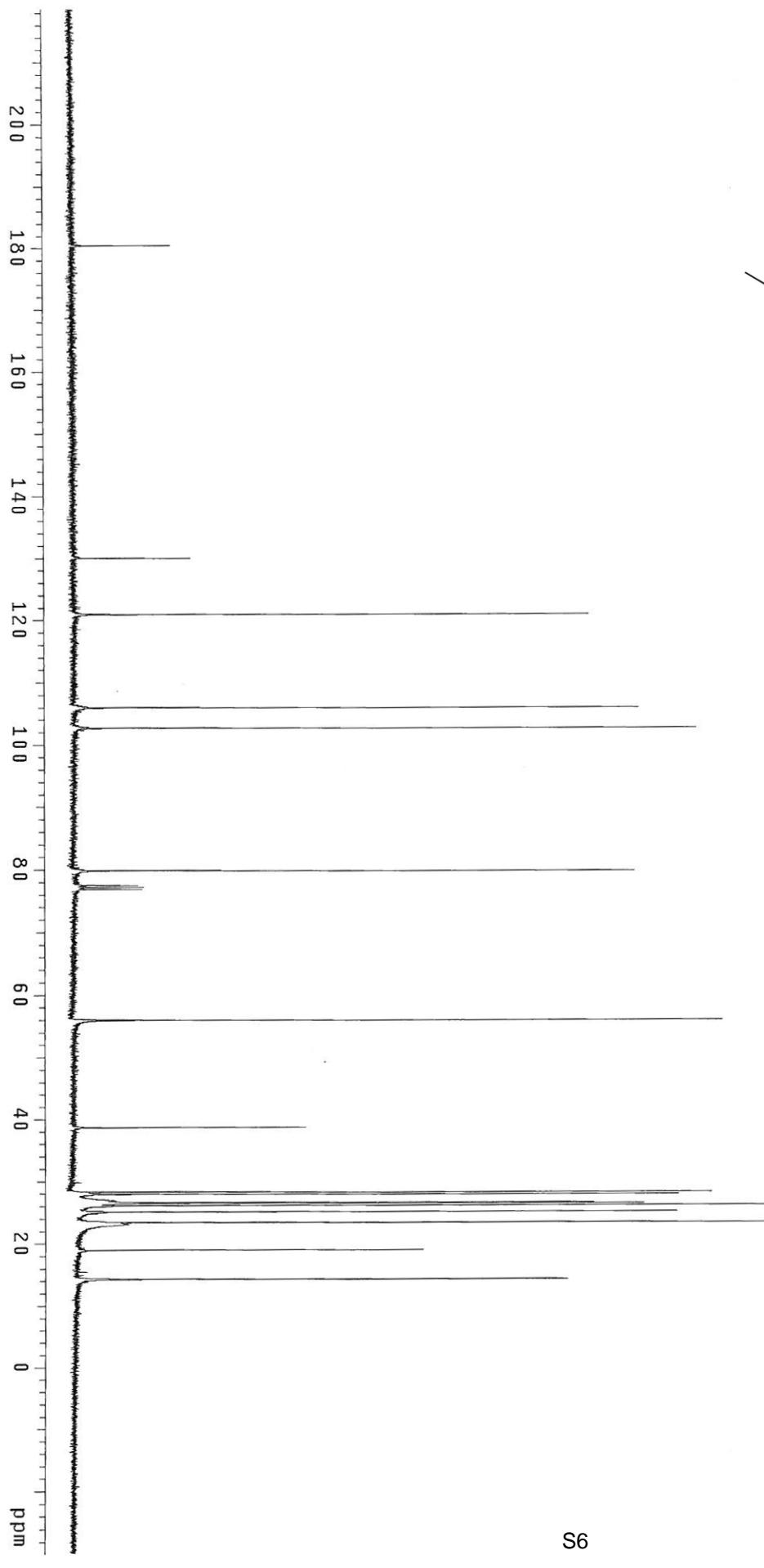
Entry	Reagent	Solvent	Temp., Time	Result ^a
1	Propylamine	Propylamine	Reflux, 1 h	1-BBu₂ (intact), 1 (none)
2	Glycine	THF/MeOH, 1:1	Reflux, 1 h	1-BBu₂ (major), 1 (minor)
3	DTT, Pyridine	THF	Reflux, 1 h	1-BBu₂ (none), 1 (none), extensive byproducts
4 ^b	TFA, 450 mM	CH ₂ Cl ₂	RT, 10 min	1 (~50%), unknown (~50%)
5 ^b	Pentanol	Pentanol	Reflux, 2 h	1-BBu₂ (trace), 1 (~quant.)
6	PhOH, K ₂ CO ₃	CH ₃ CN	Reflux, 2 h	1-BBu₂ (trace), 1 (~quant.)
7	N ₂ H ₄ ·H ₂ O	THF	Reflux, 1 h	1-BBu₂ (none), 1 (~quant.)
8	KOH (excess)	THF/MeOH, 1:1	Reflux, 1 h	1-BBu₂ (none), 1 (~quant.)

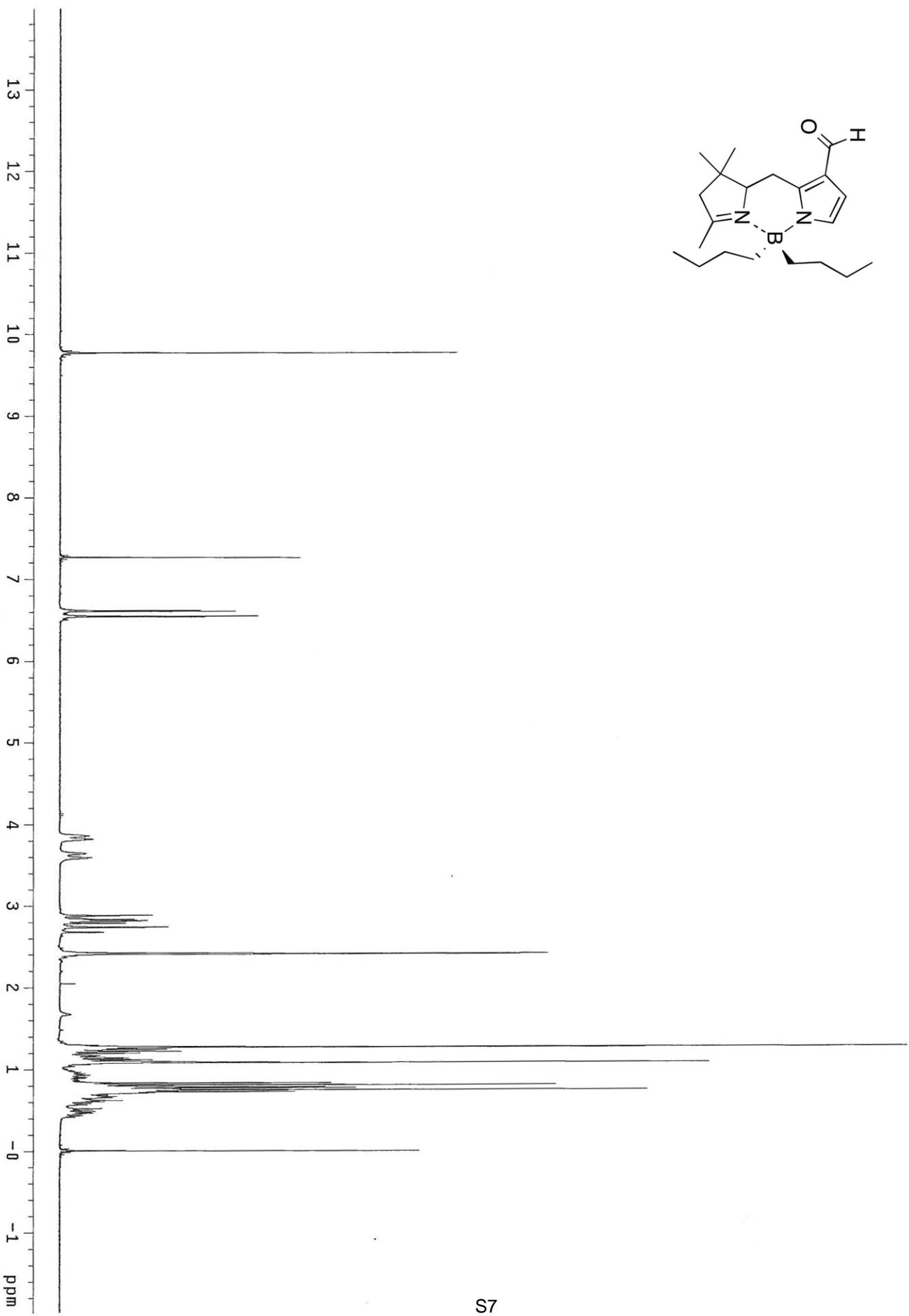
^aThe product distribution was estimated on the basis of TLC analysis with visual inspection.

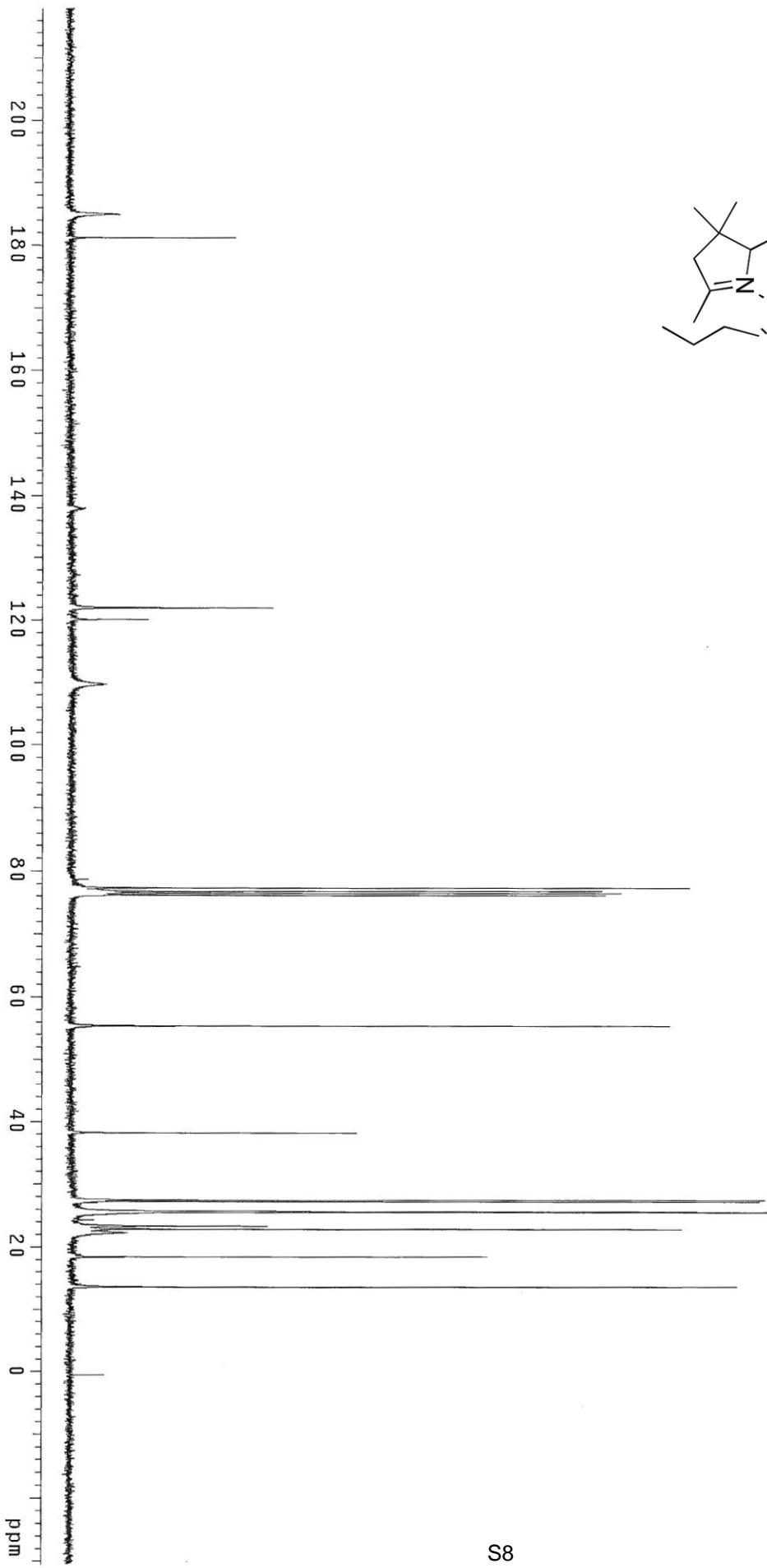
^bApplication of the same conditions to **1-BMe₂** afforded the corresponding product distribution.

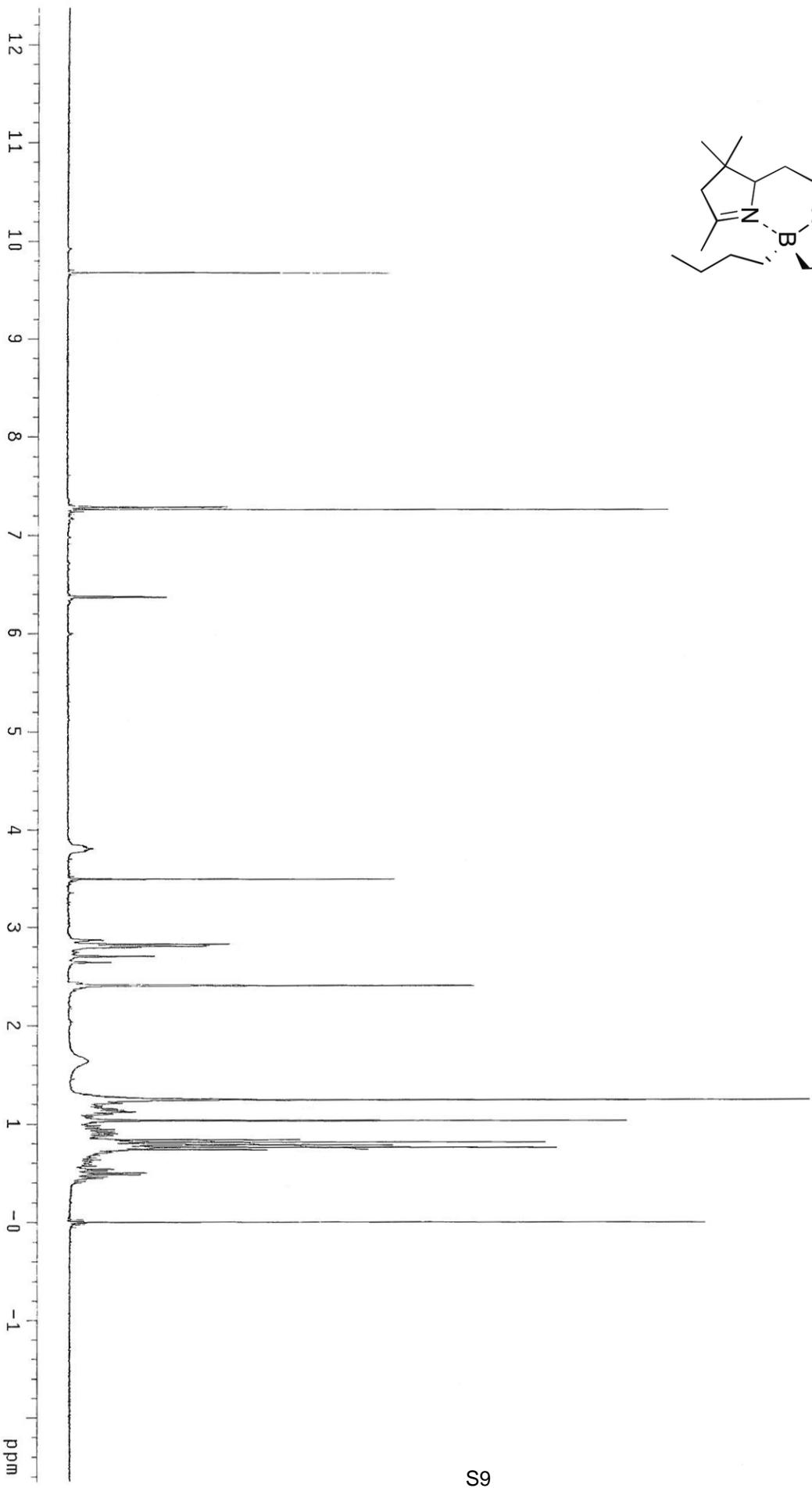
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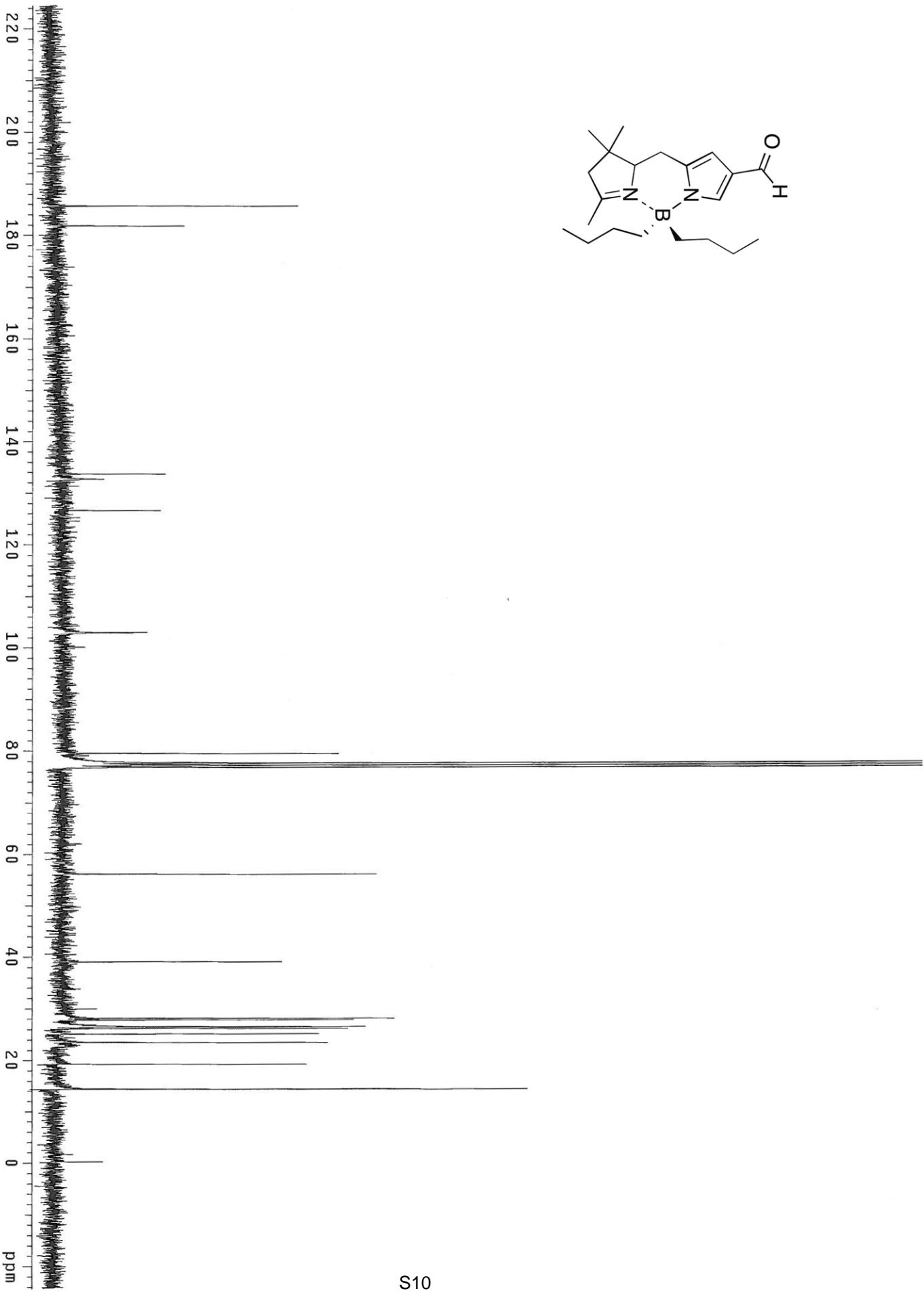


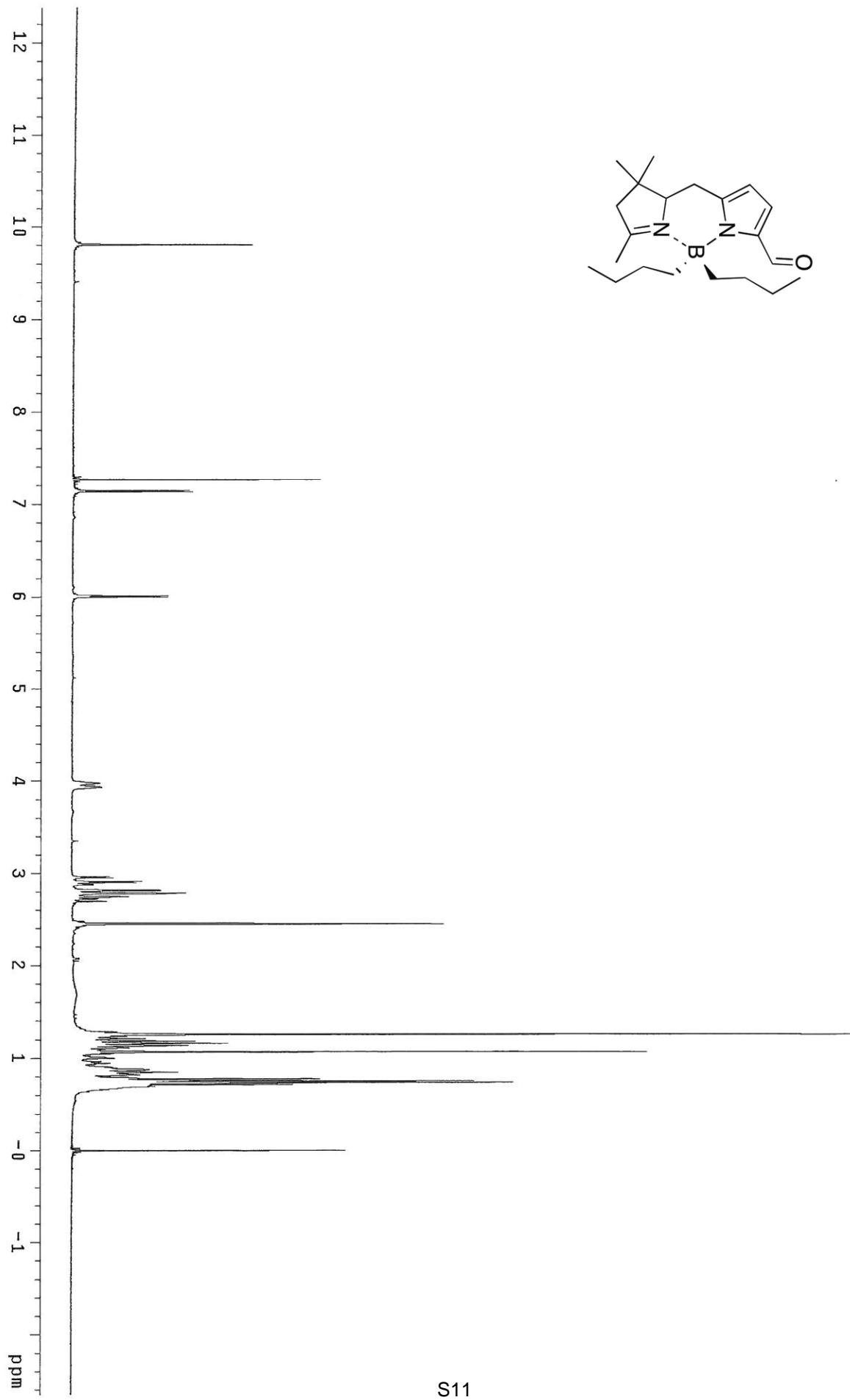


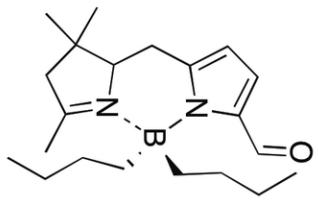
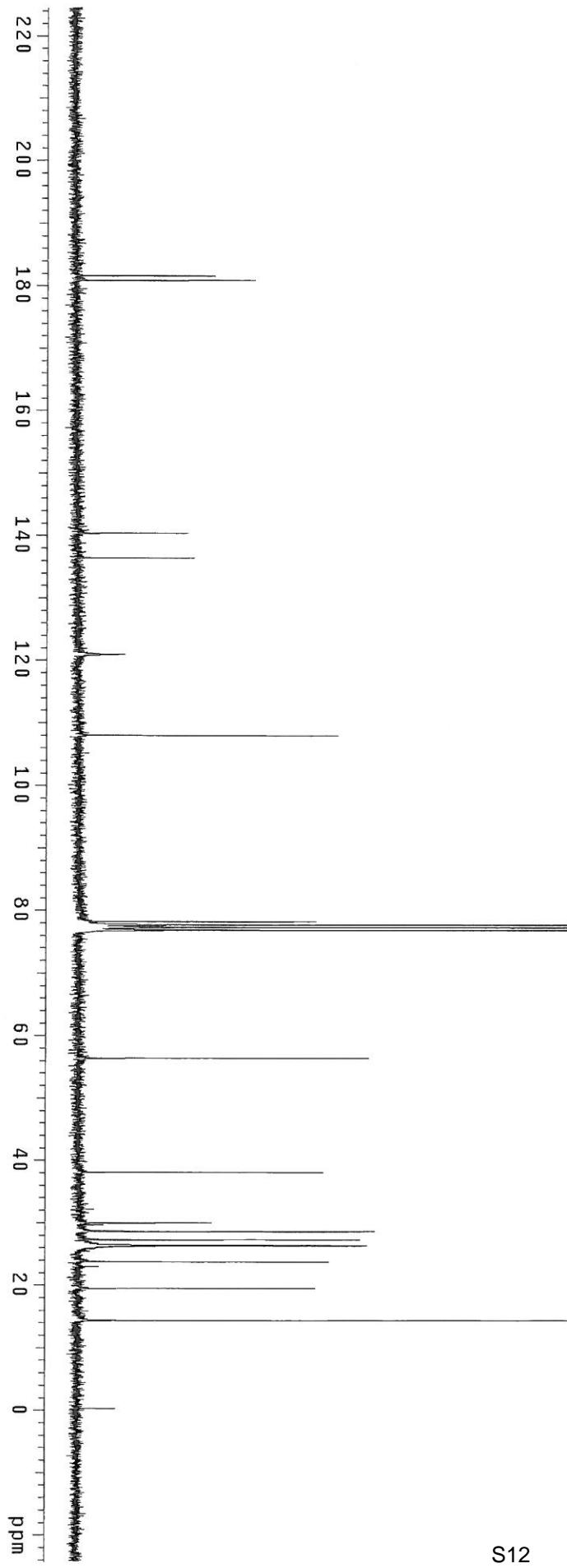


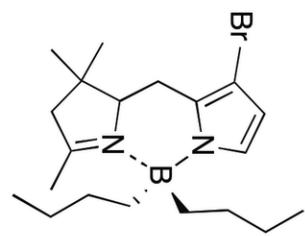
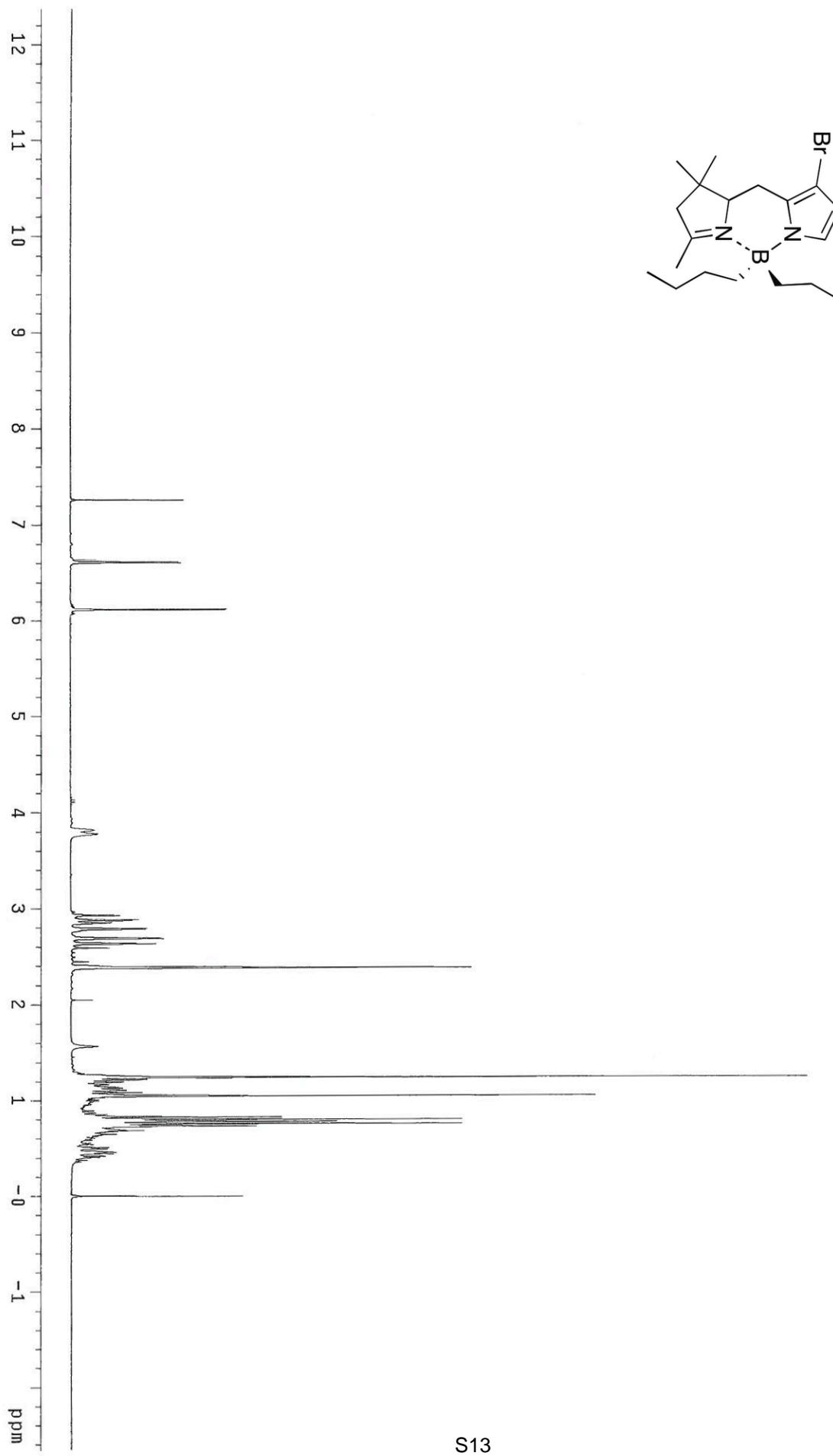


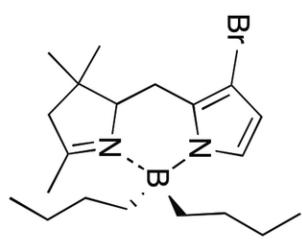
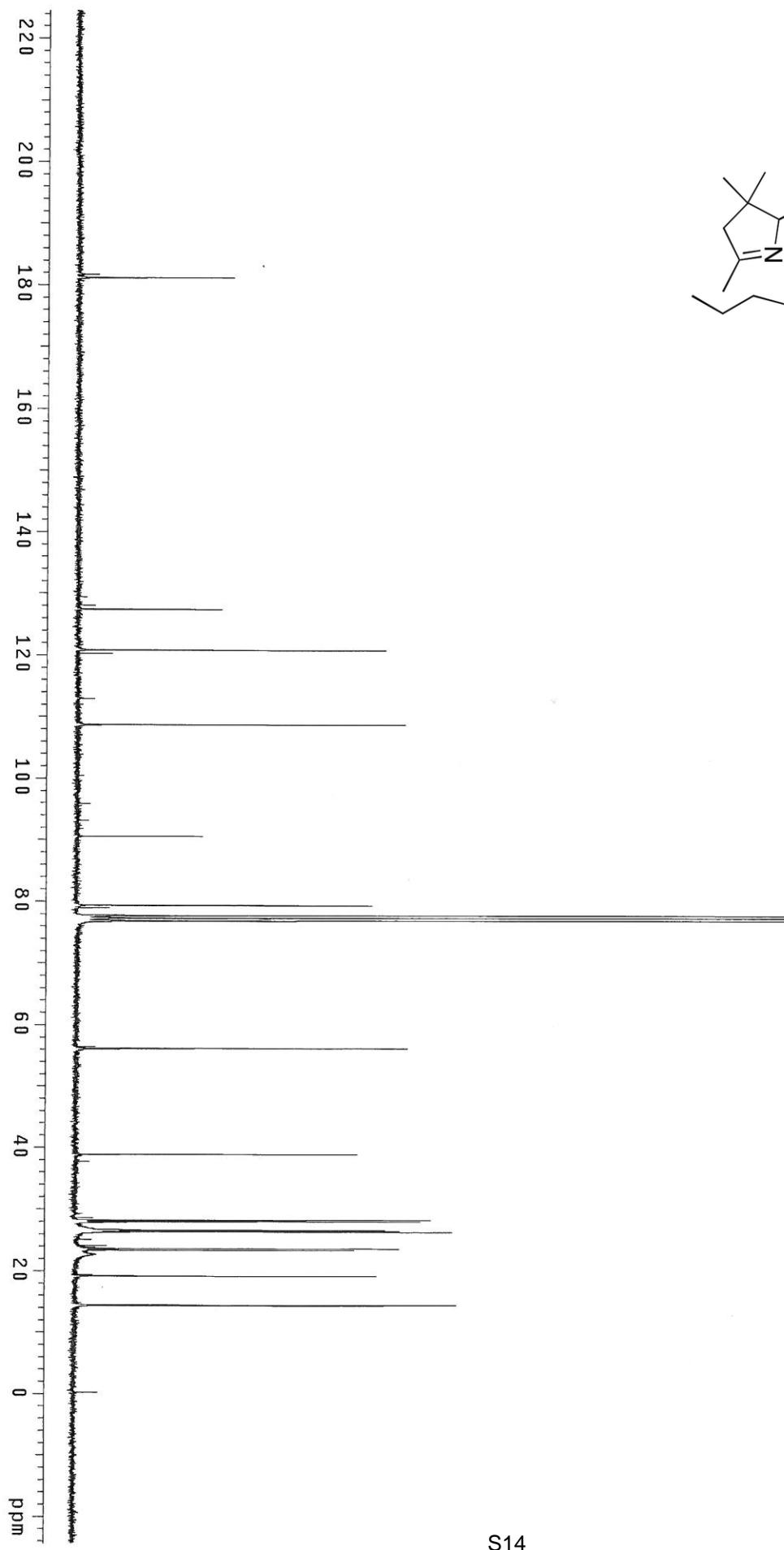


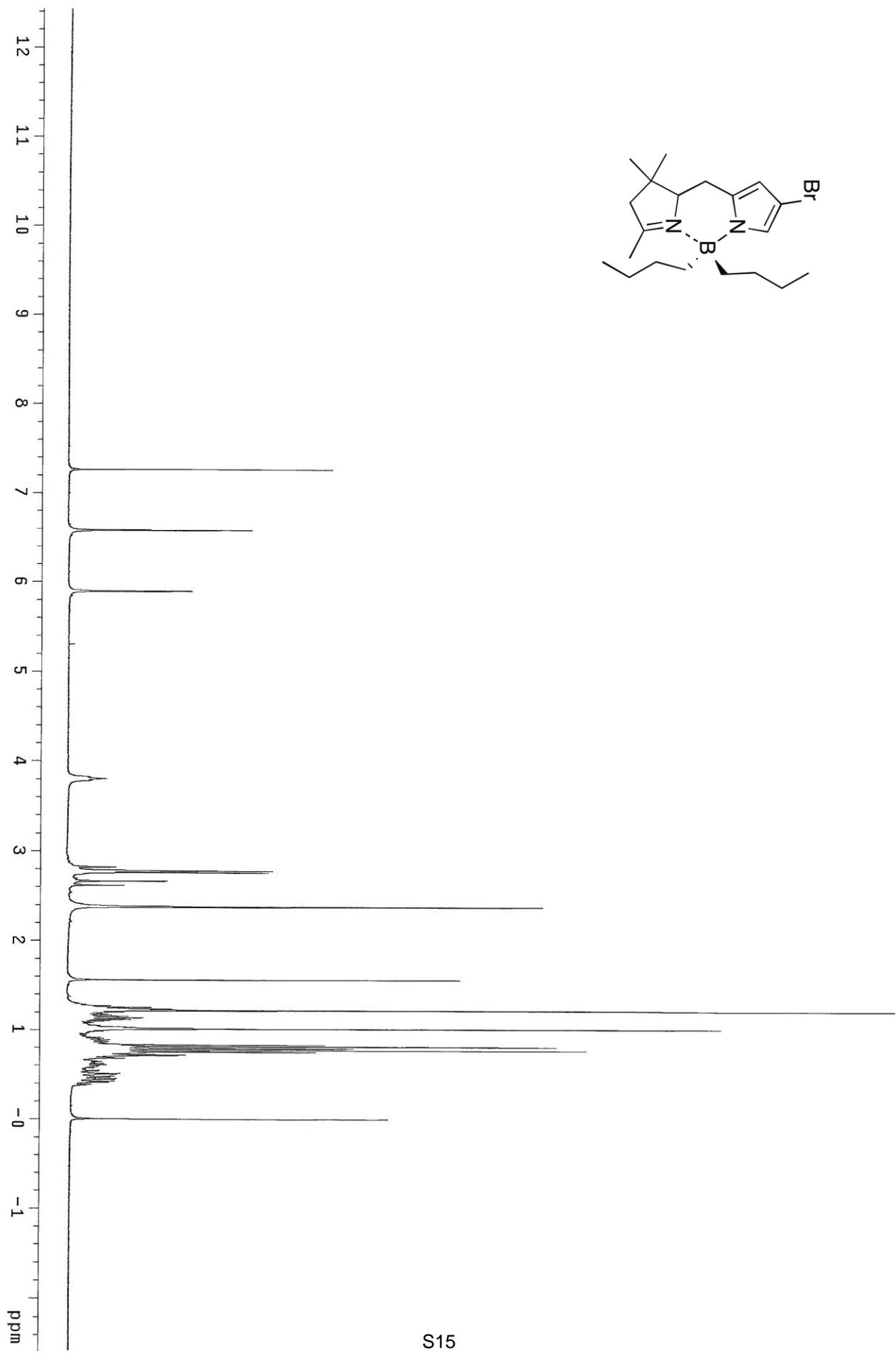


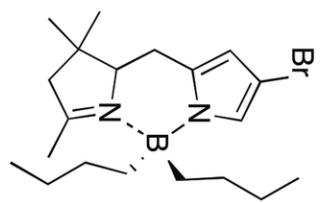
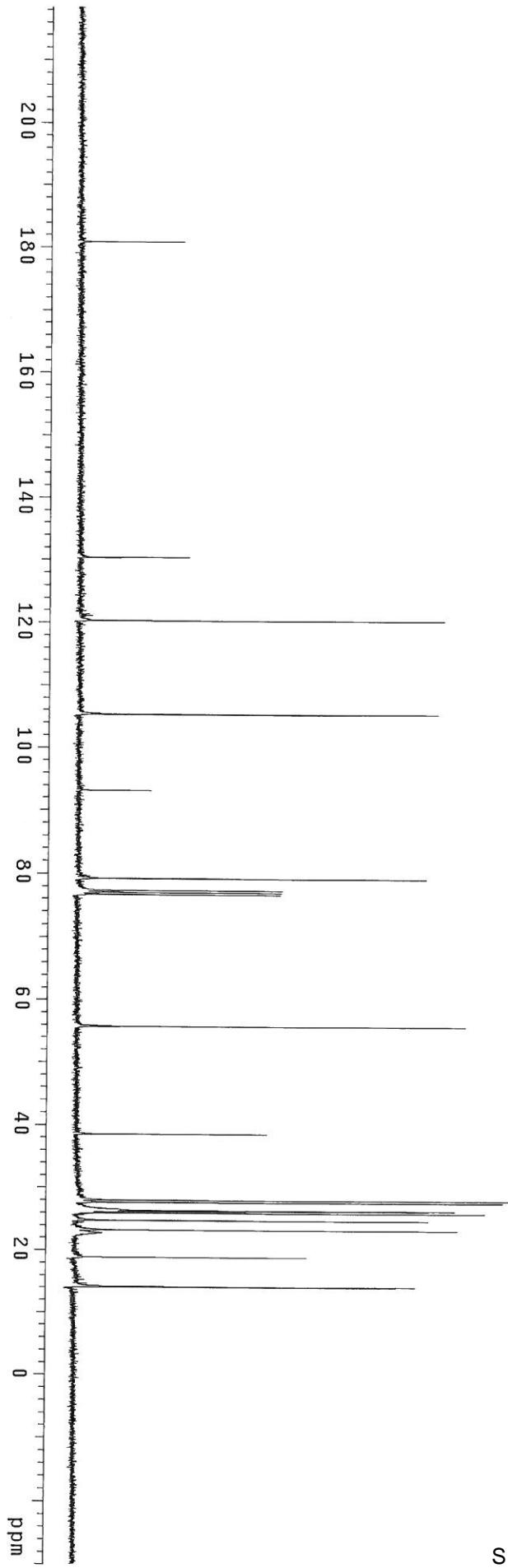


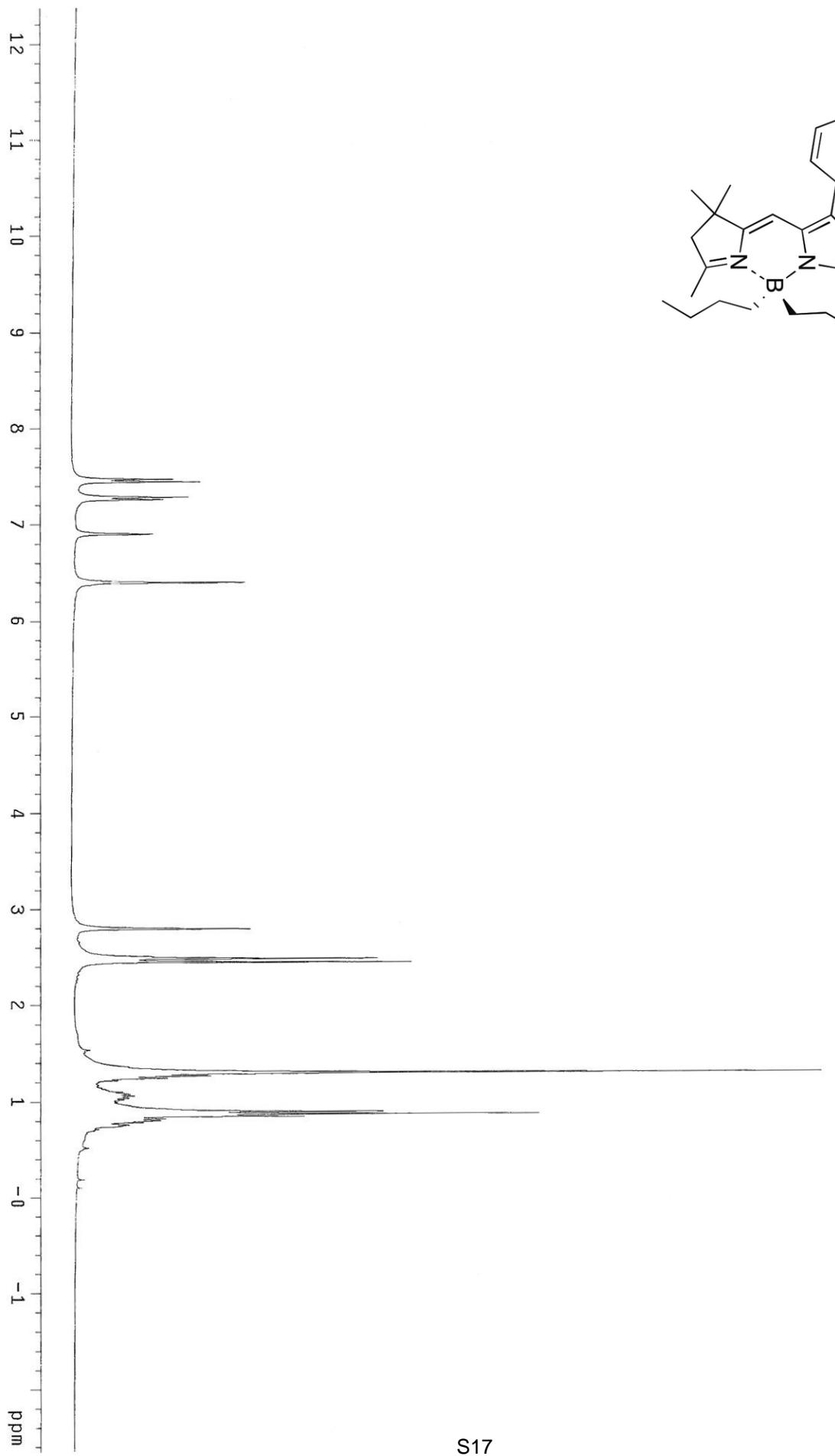


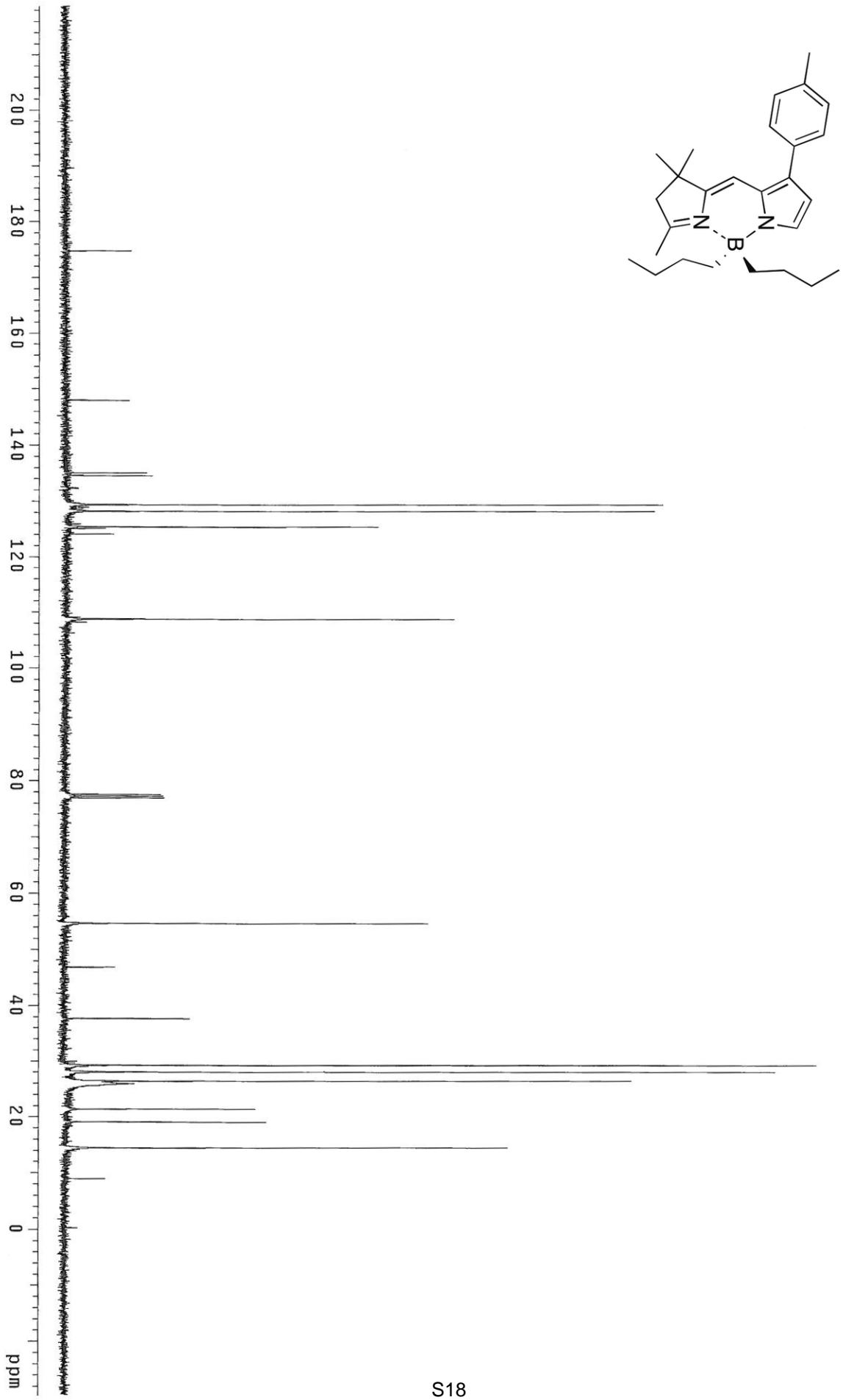


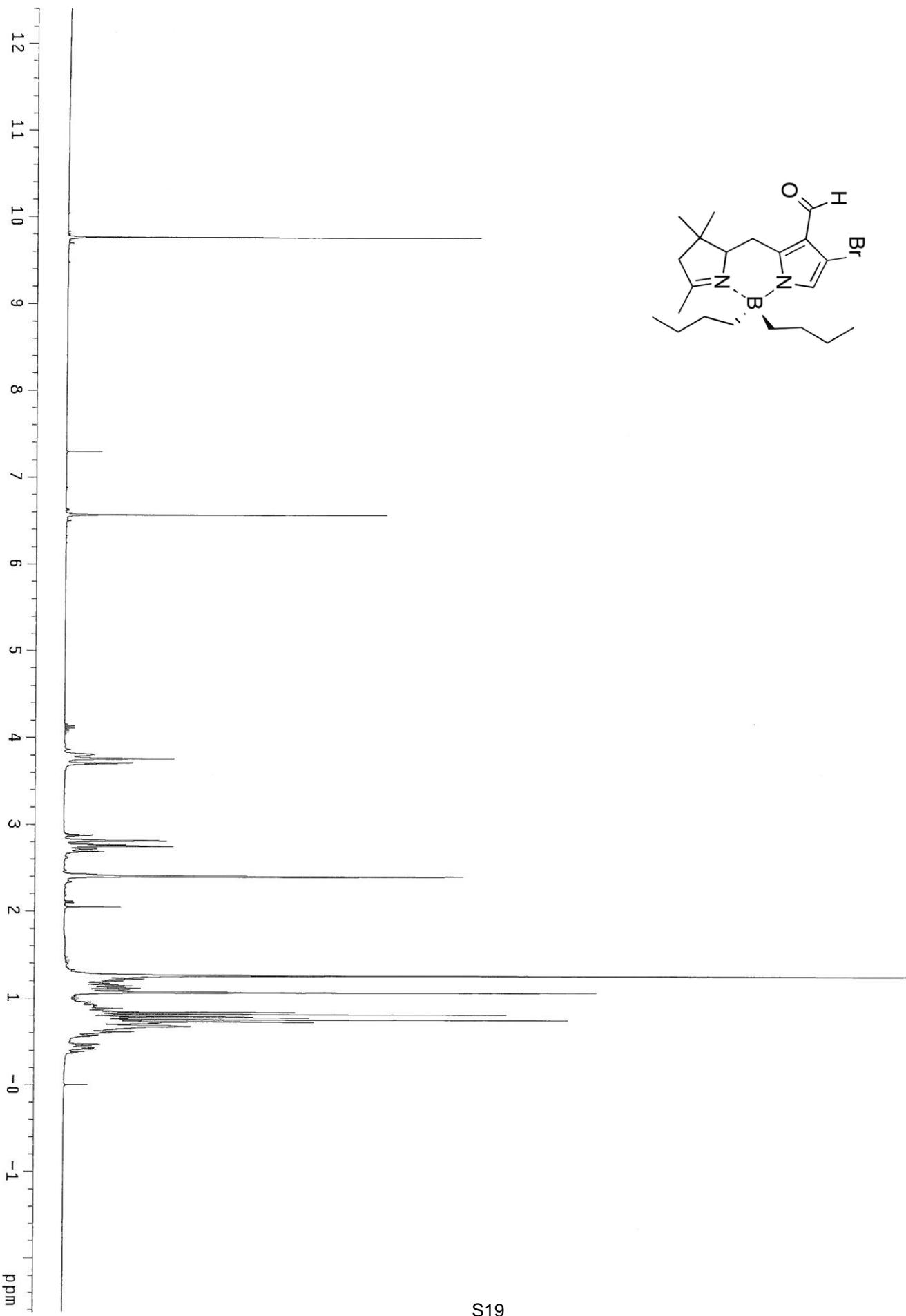


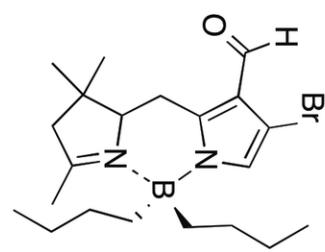
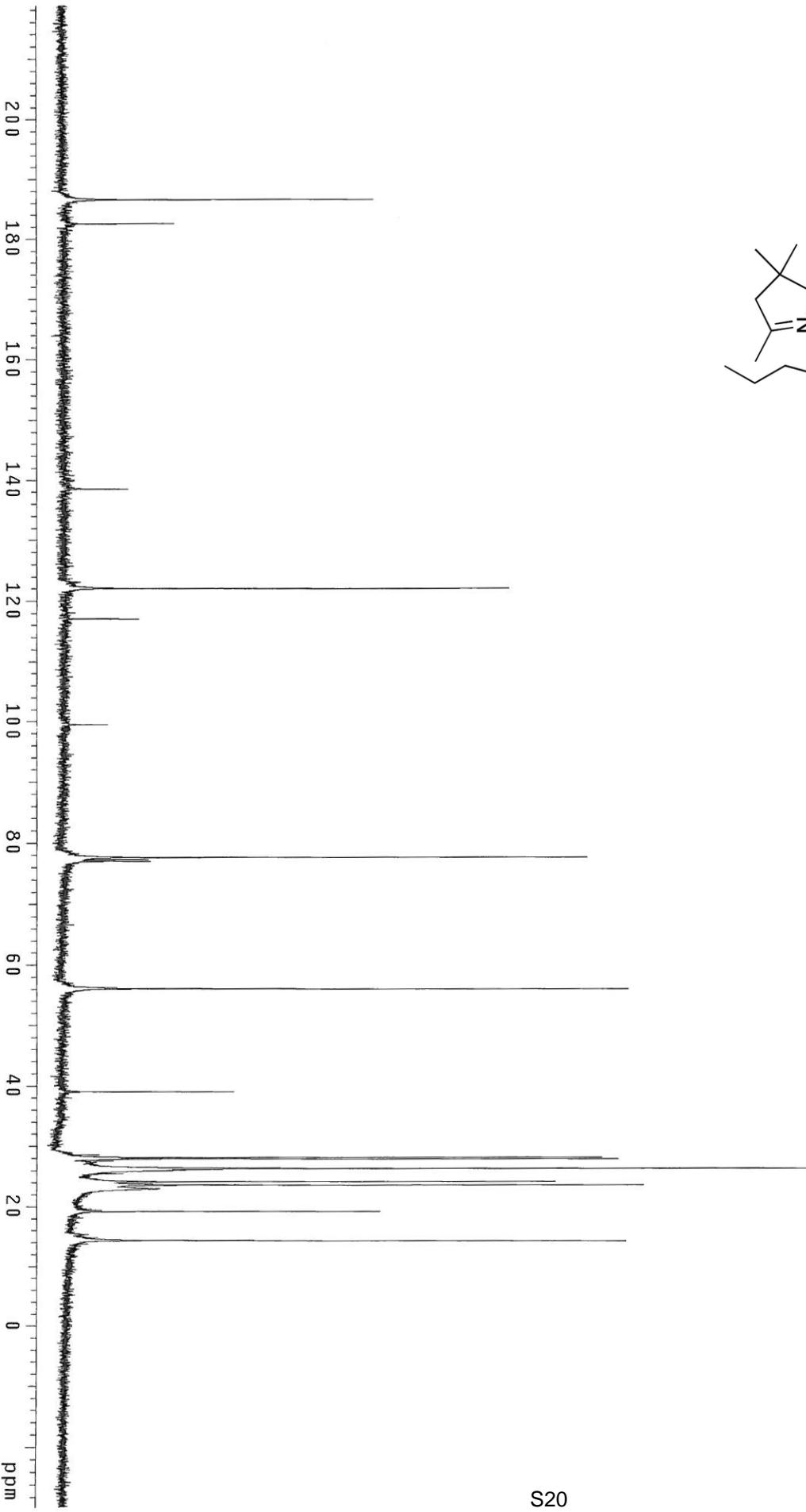


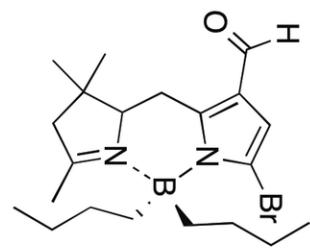
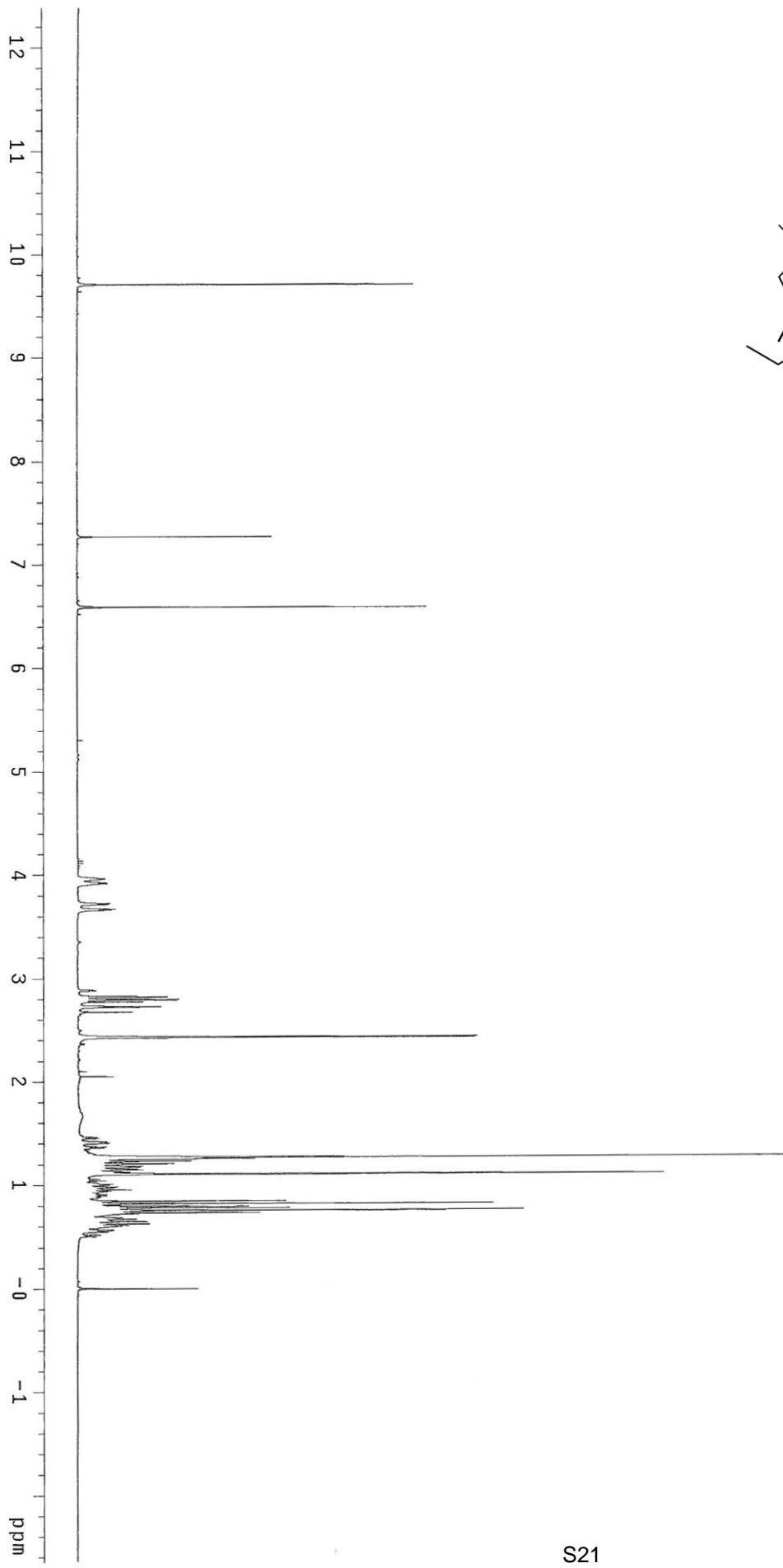


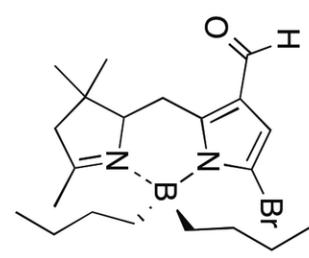
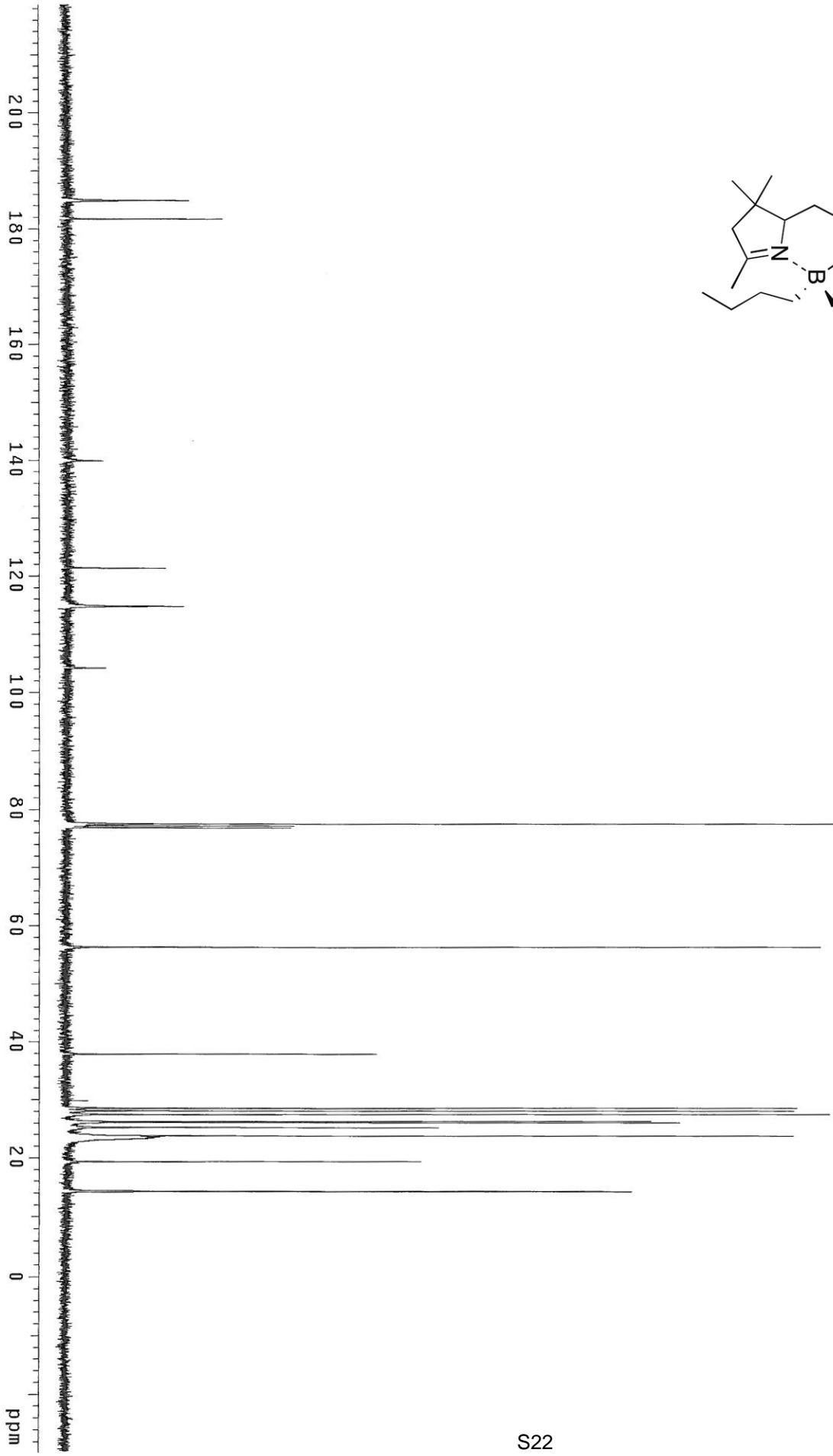


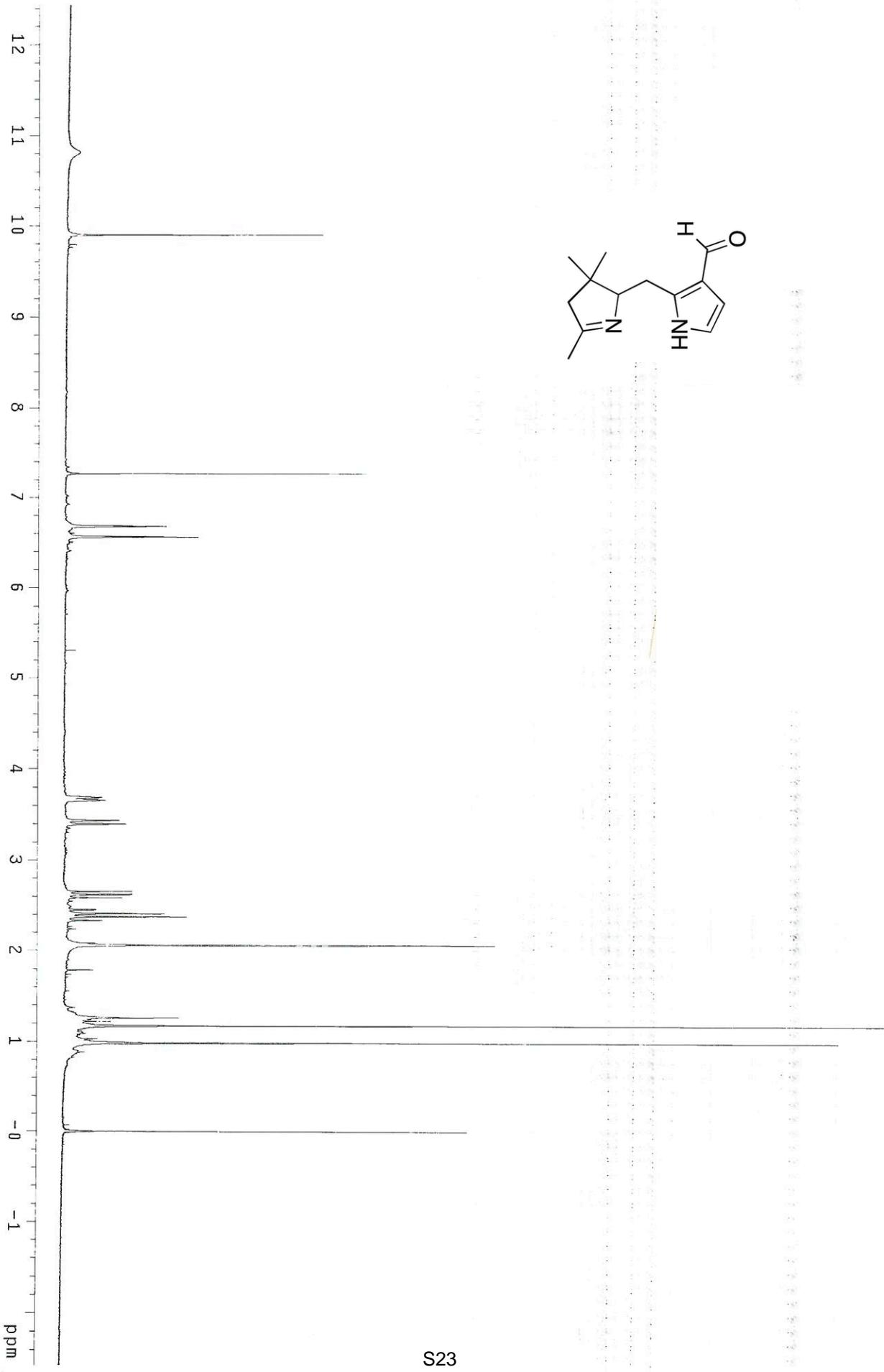


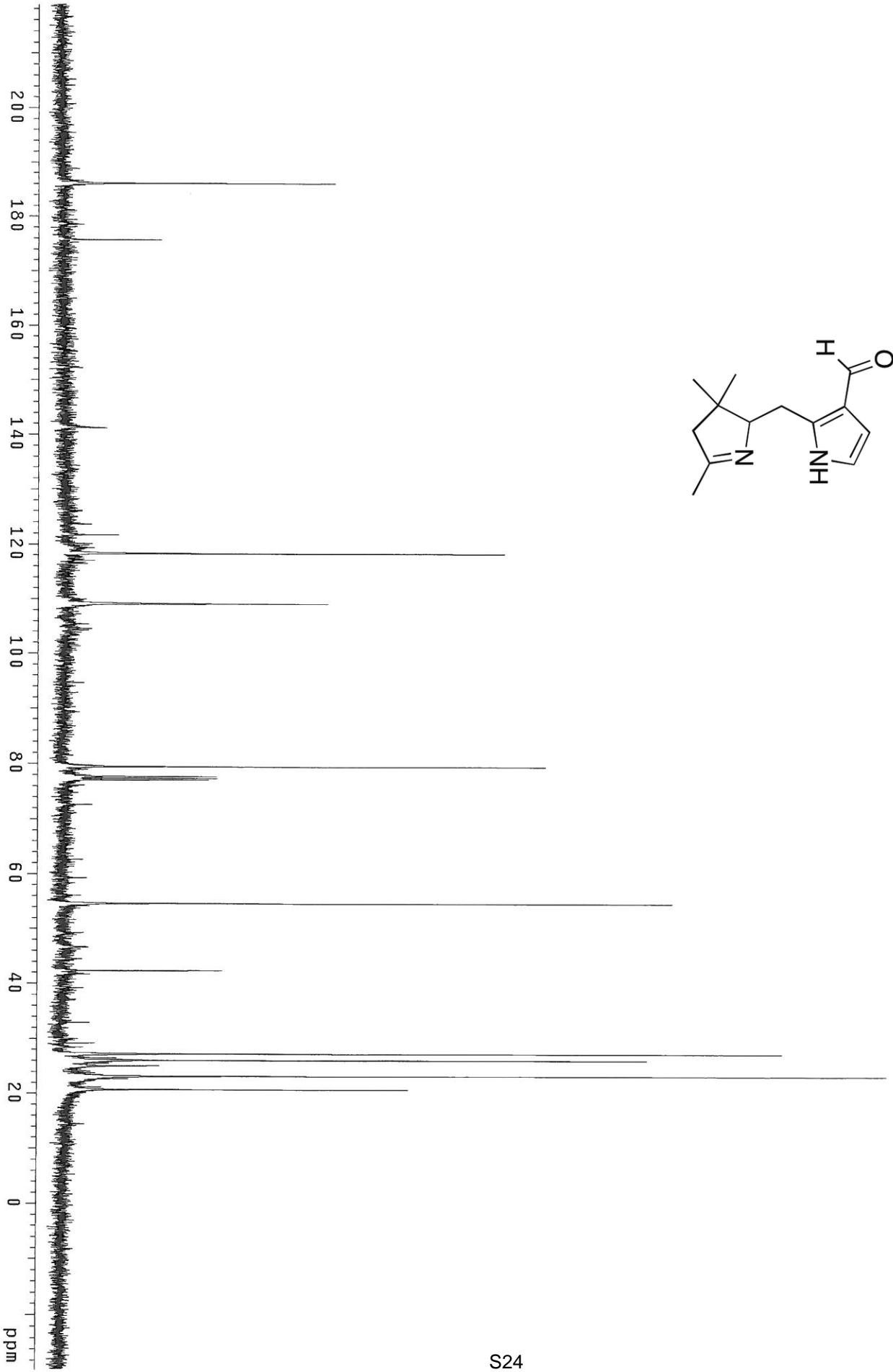


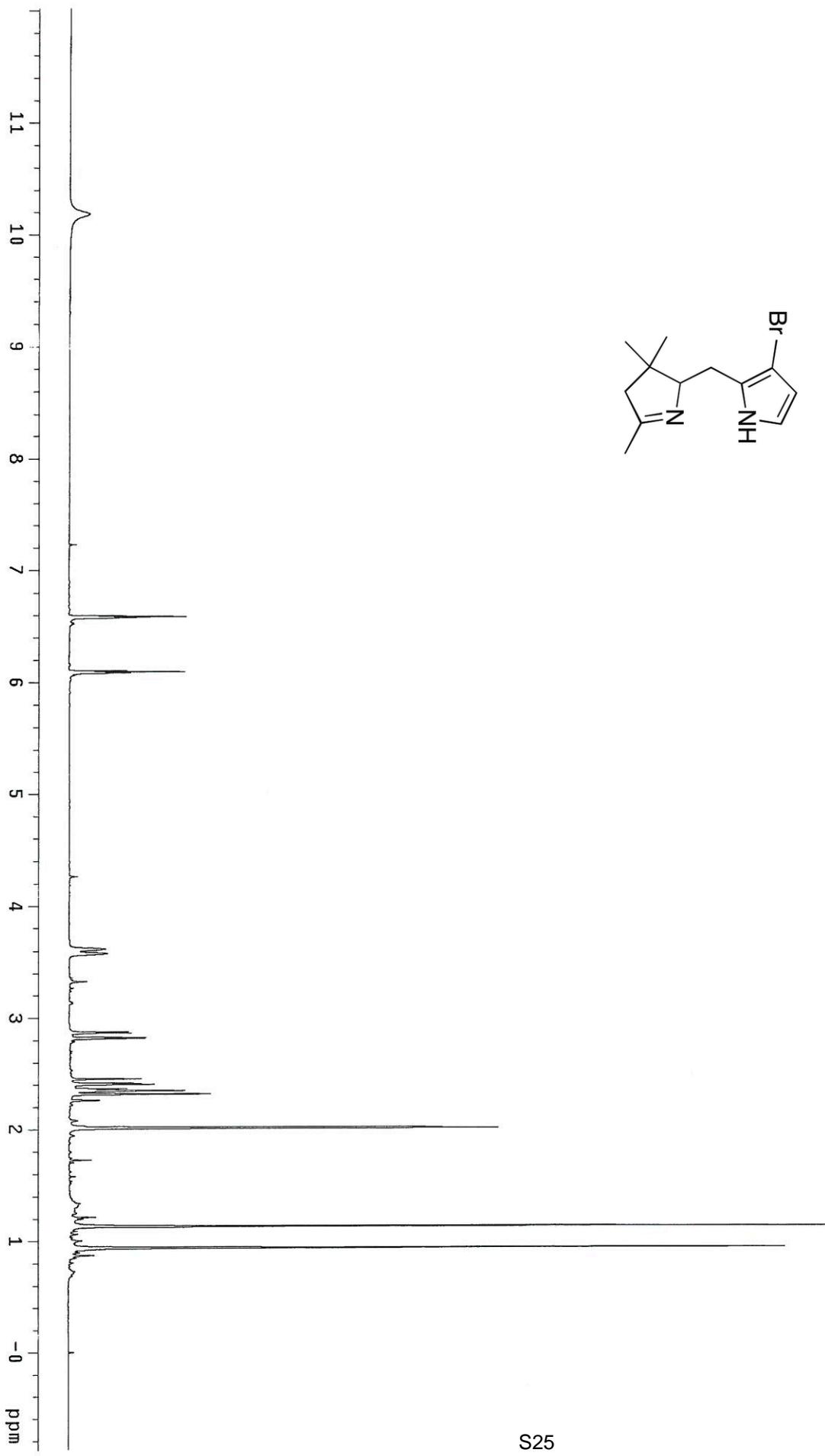


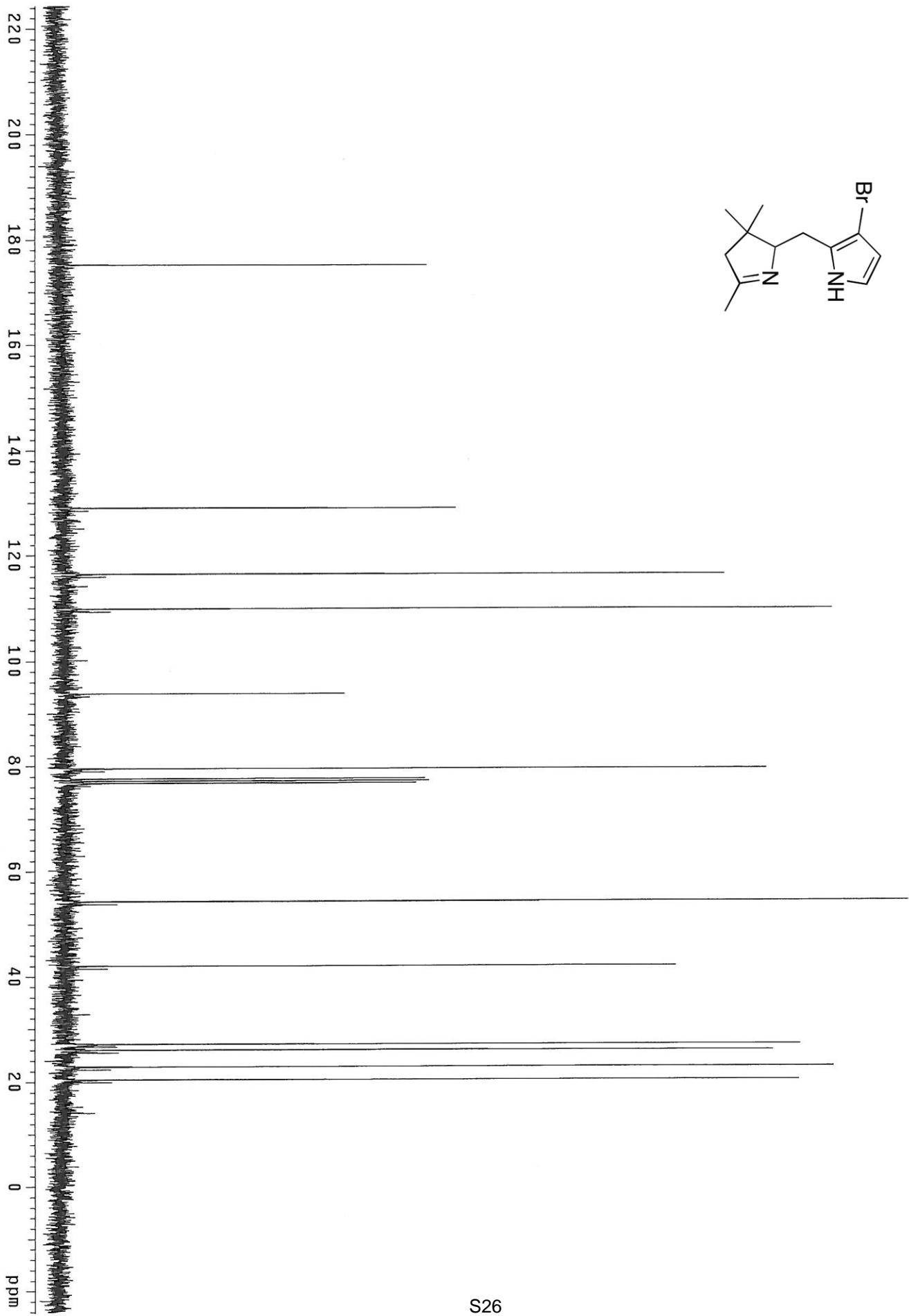


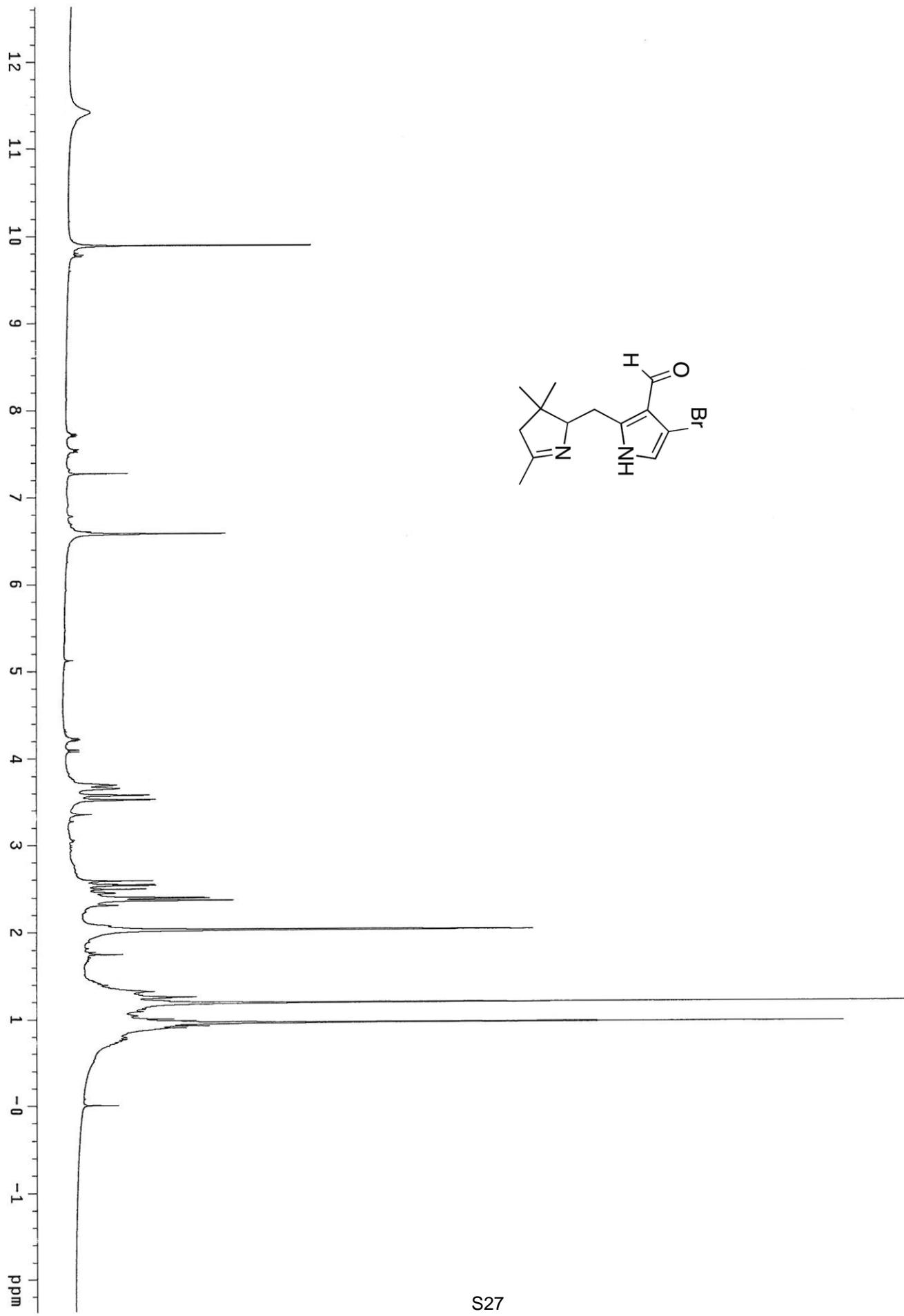


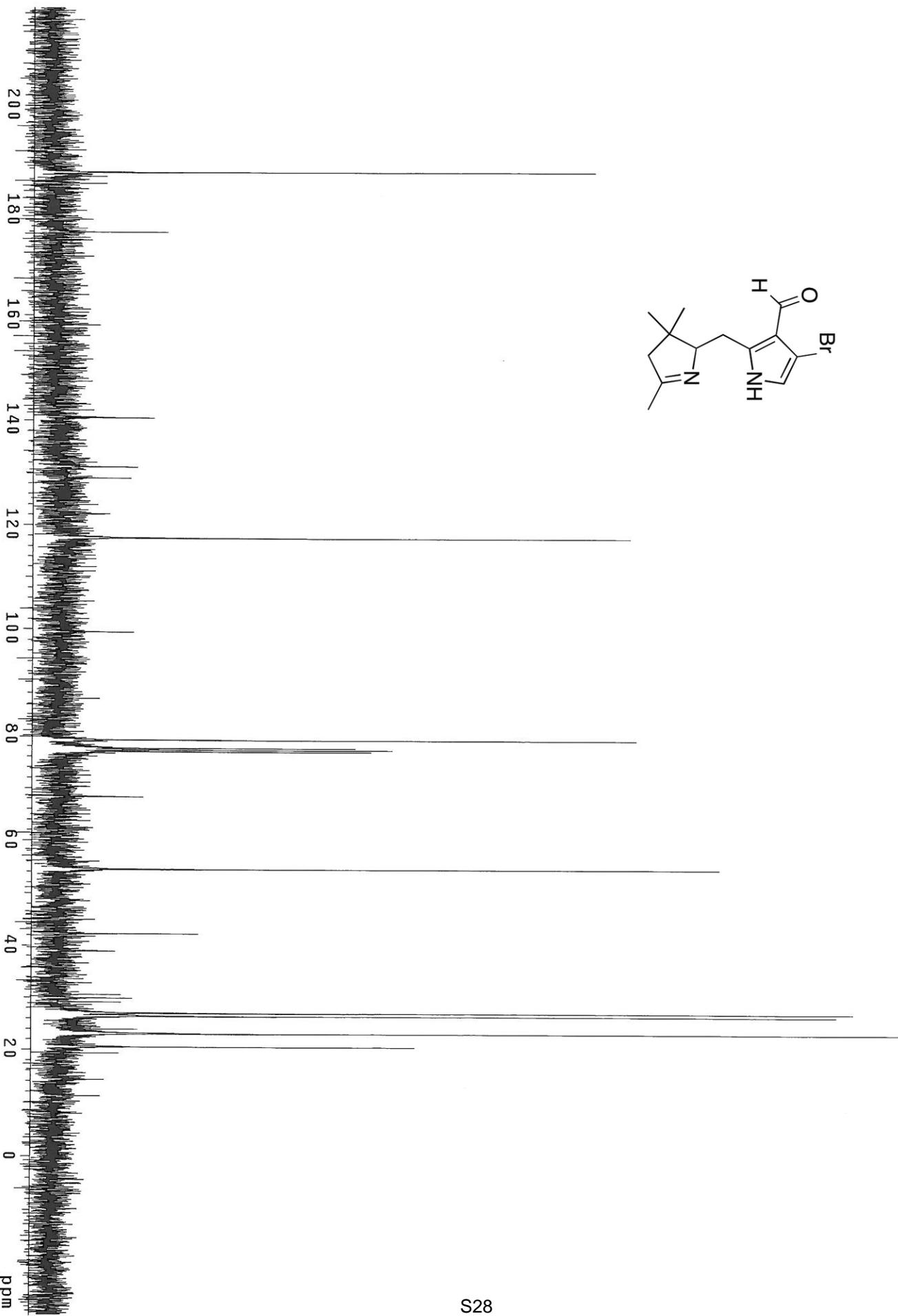


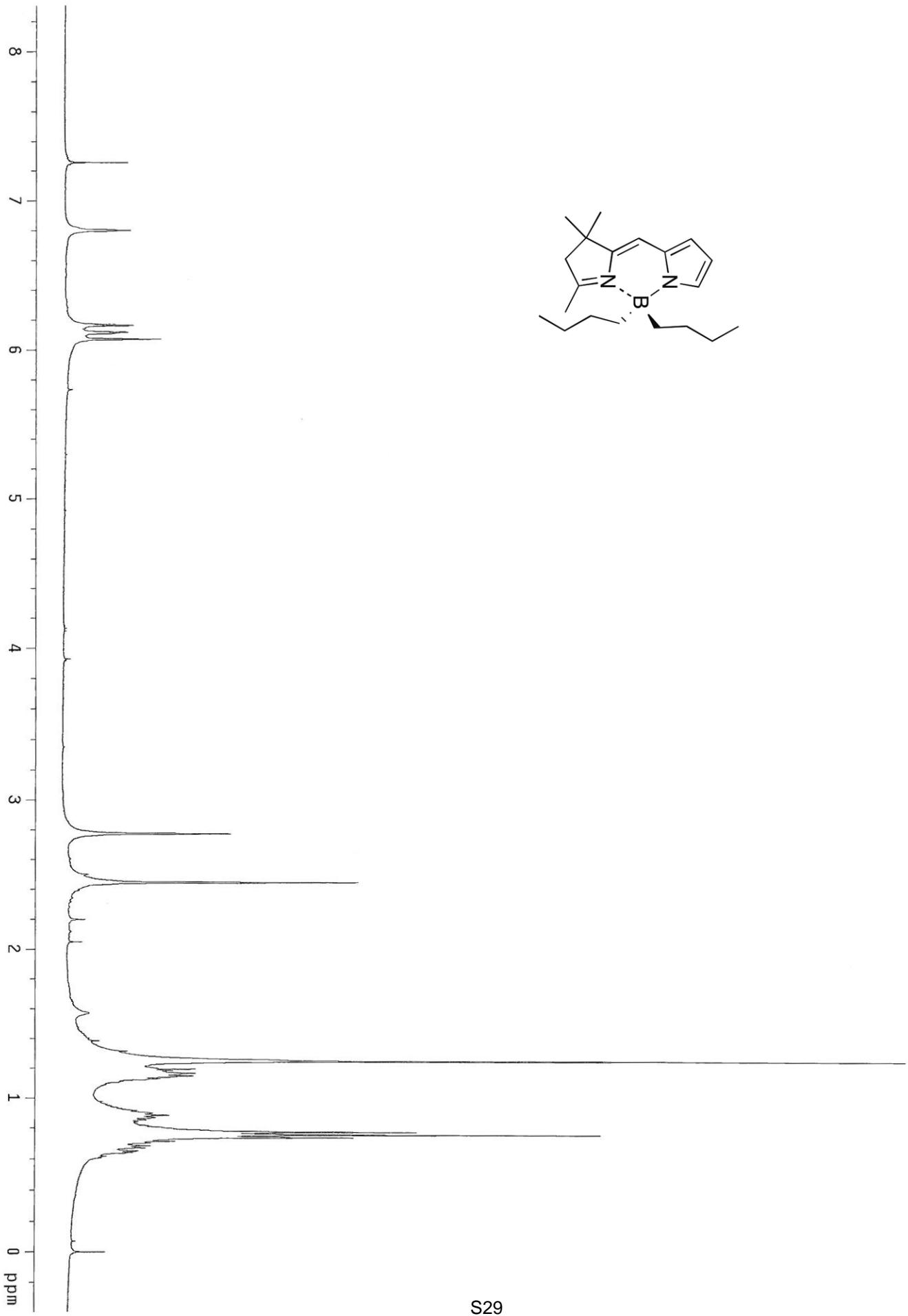


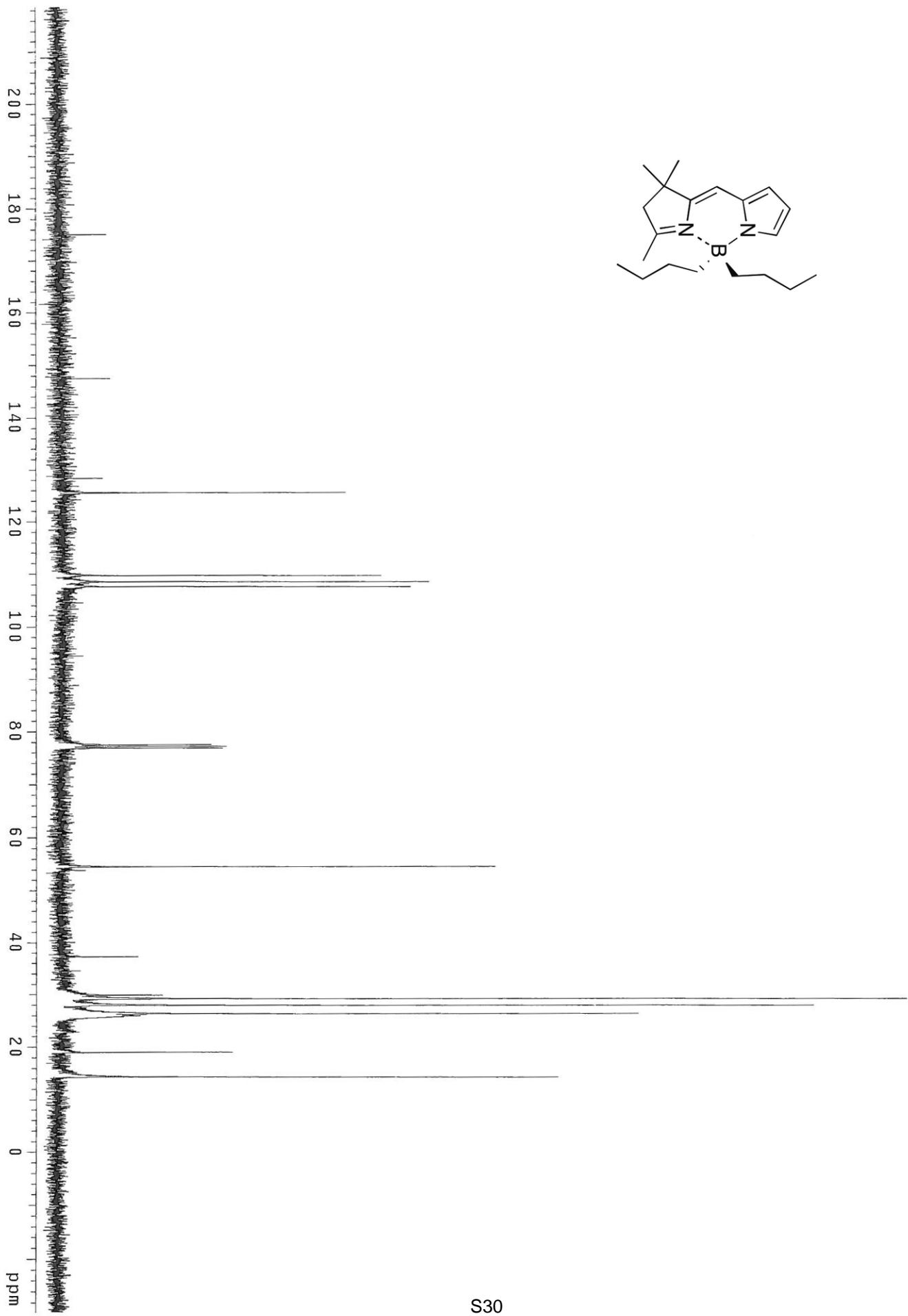


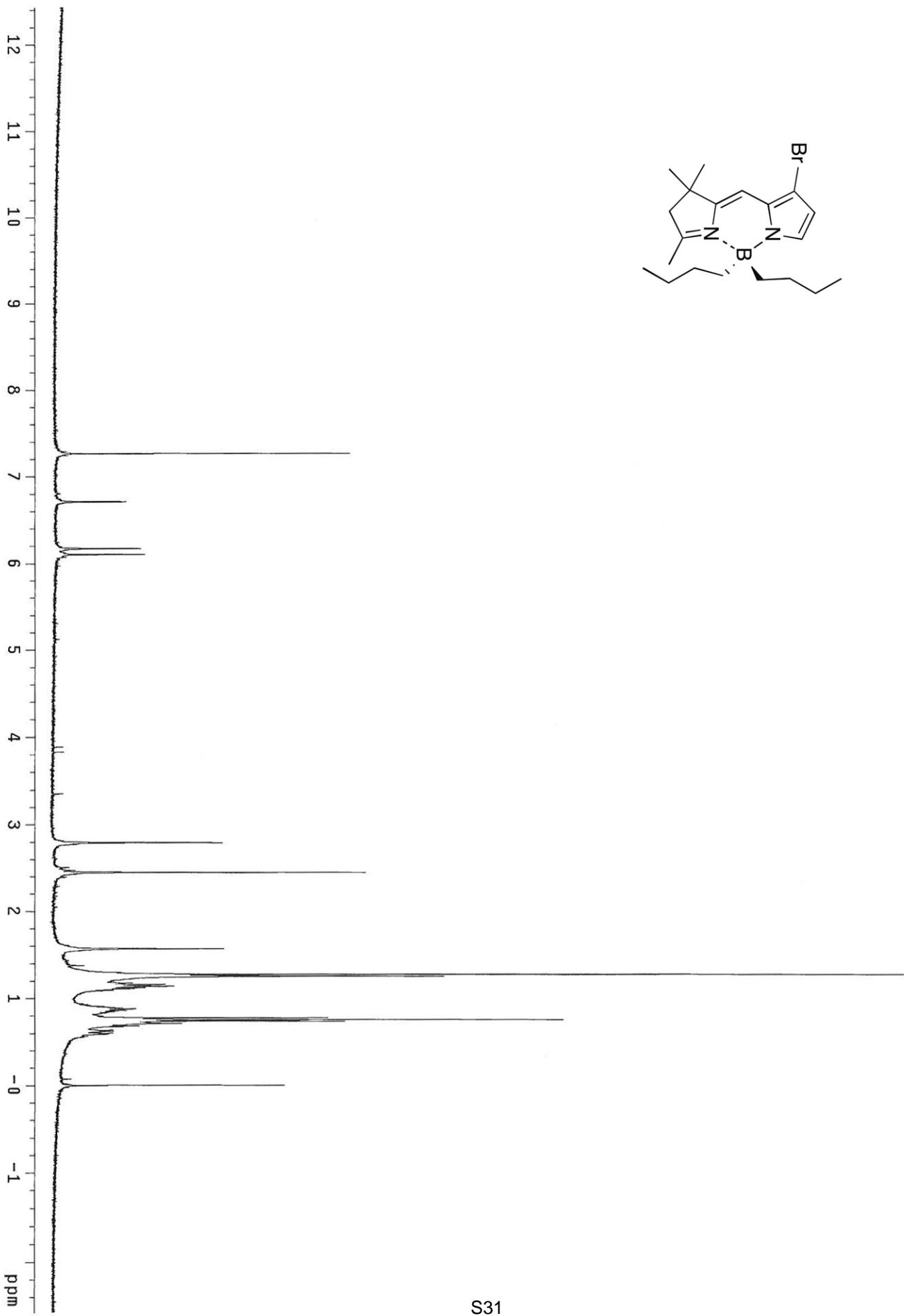


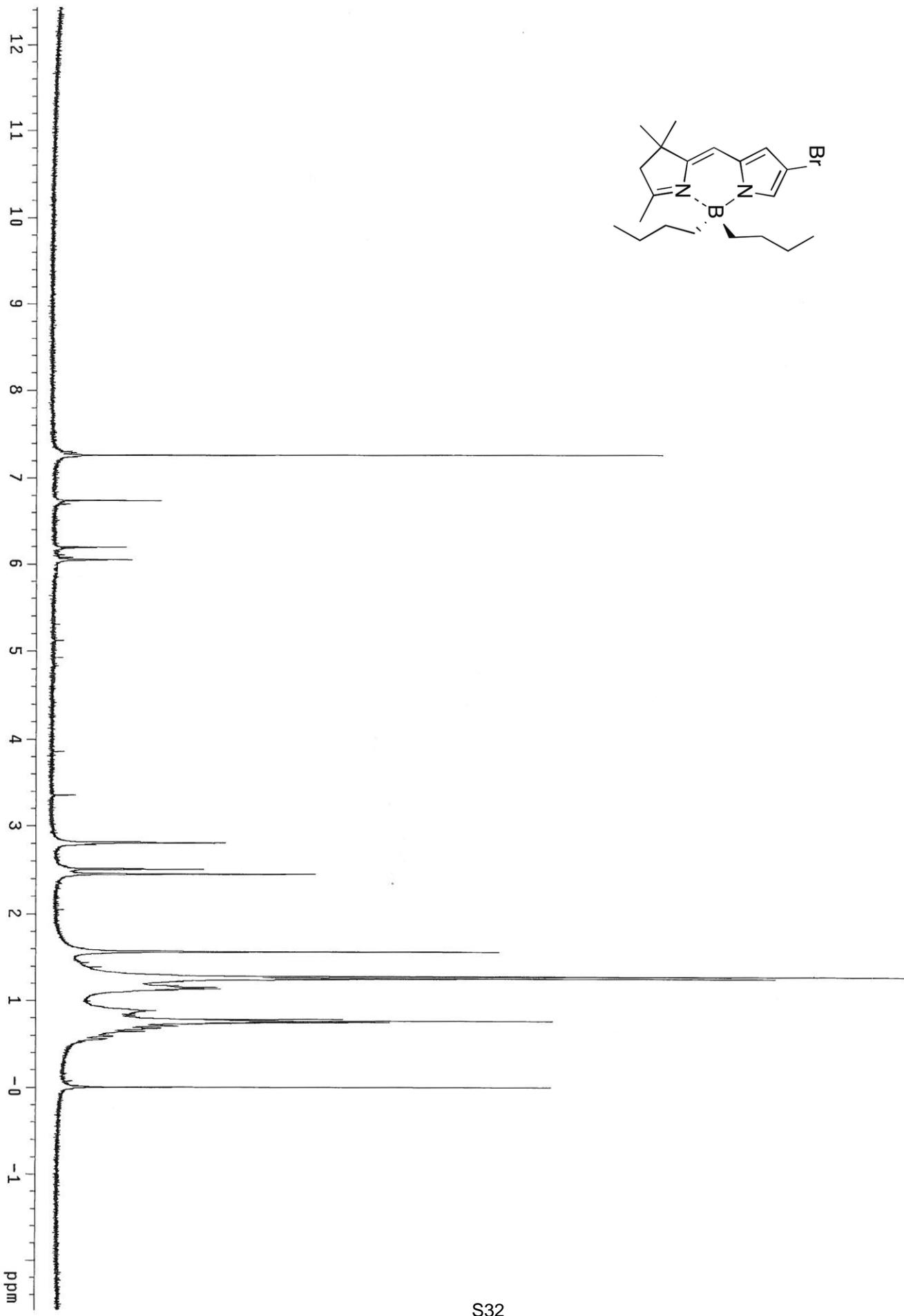


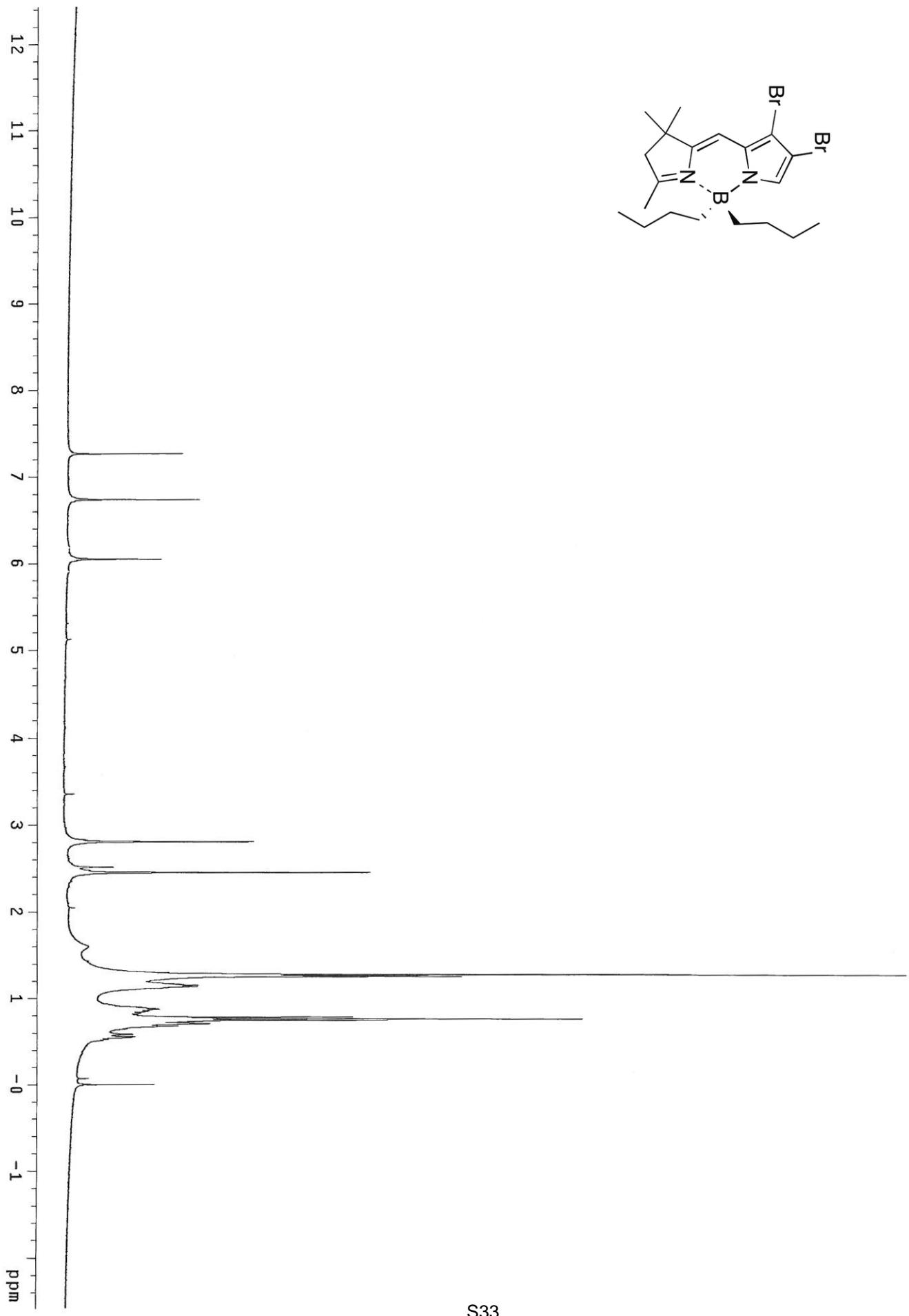


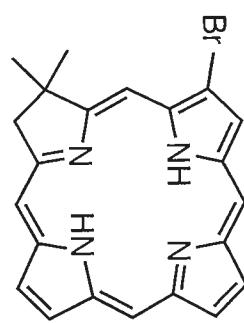
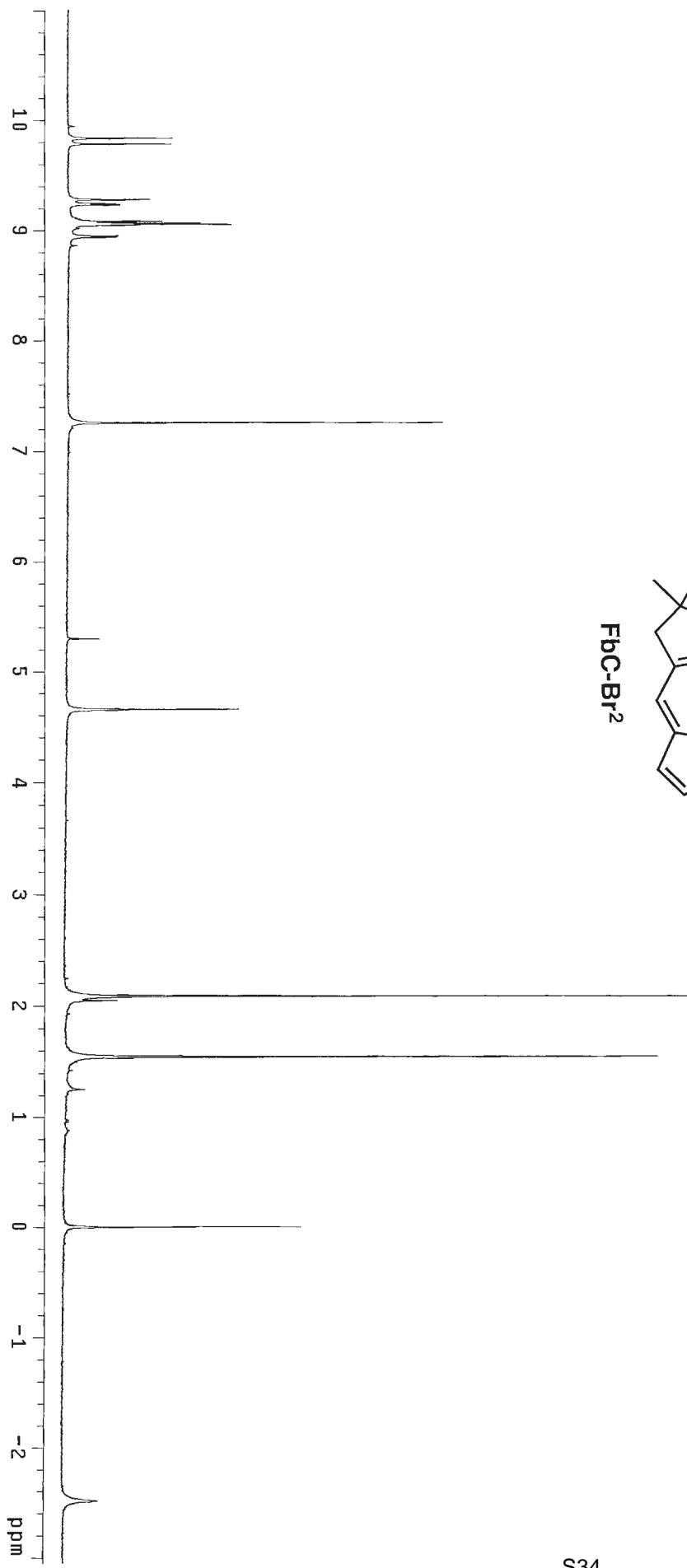




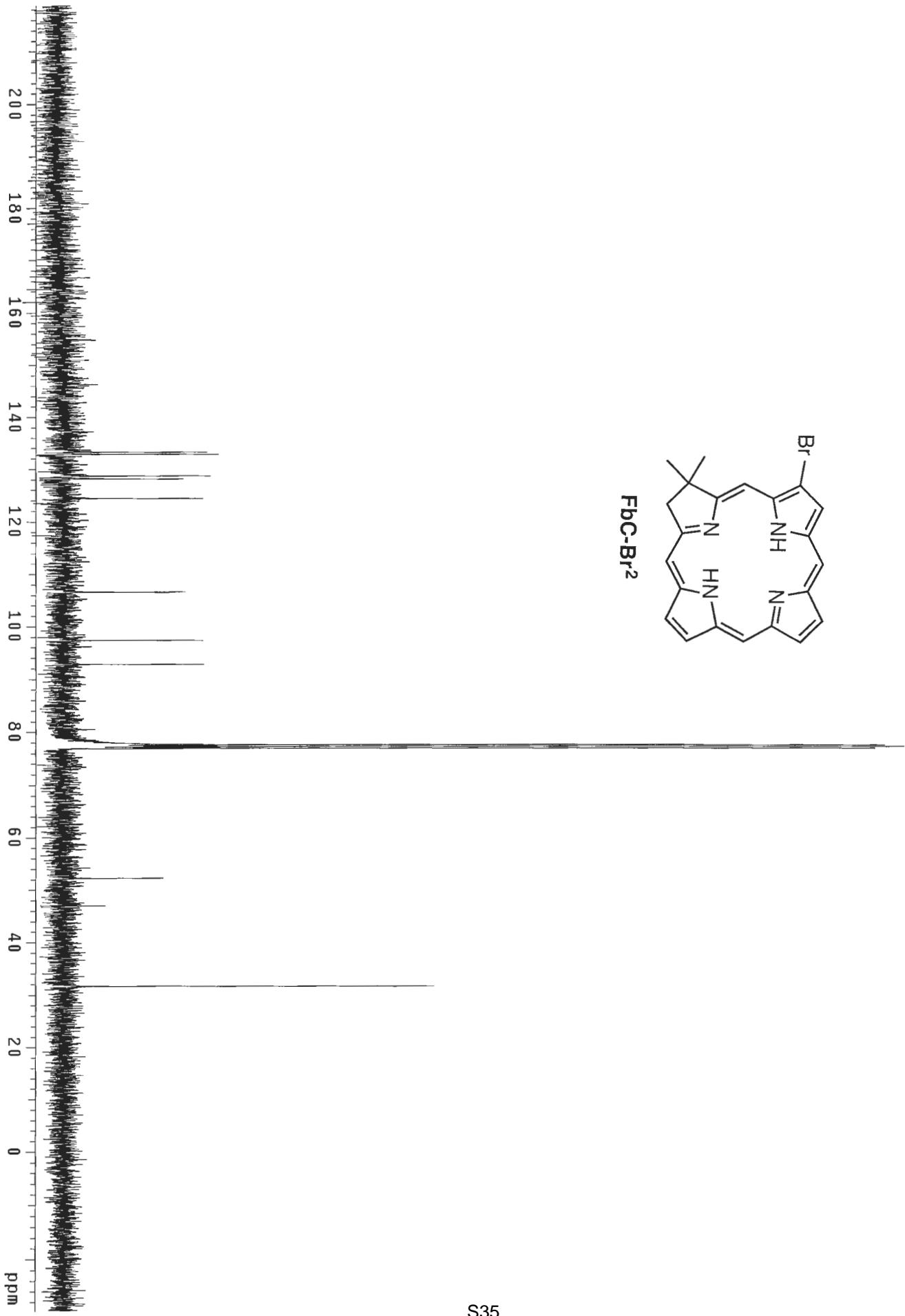






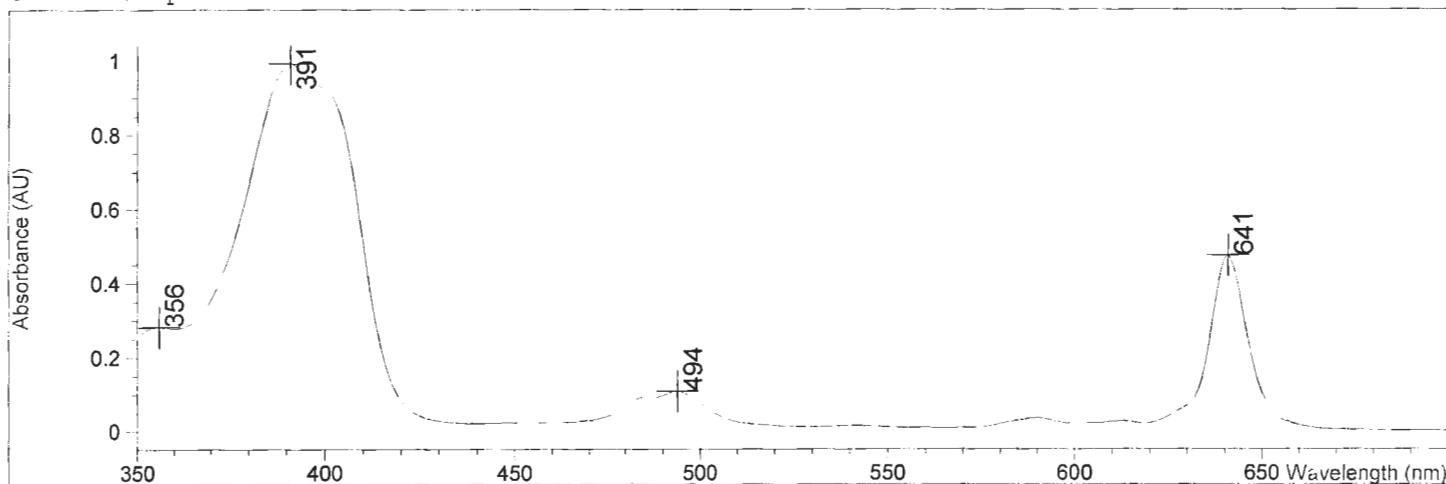


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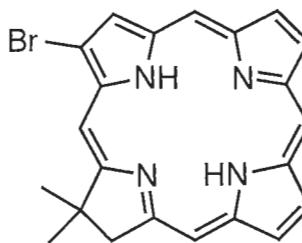


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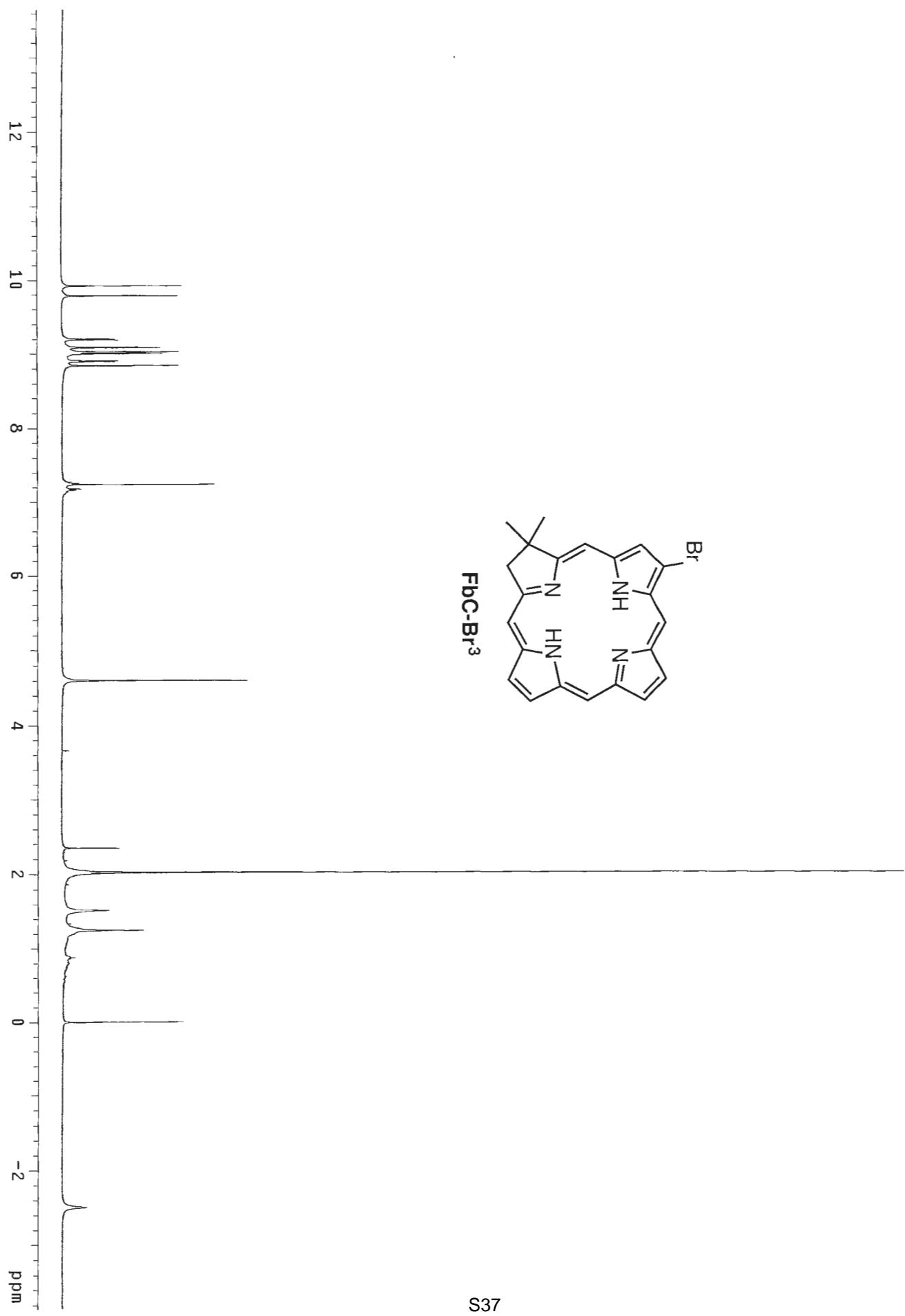
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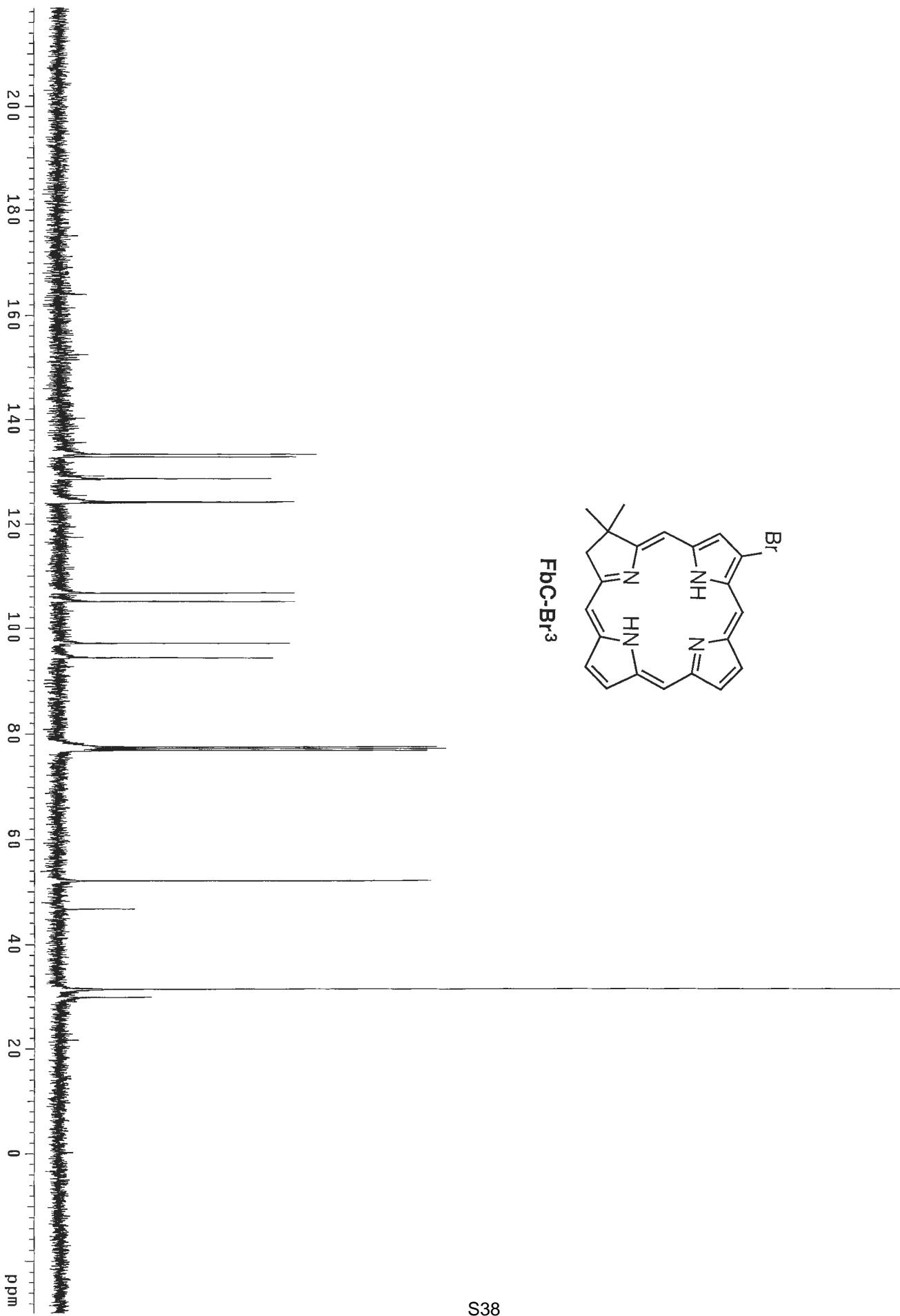


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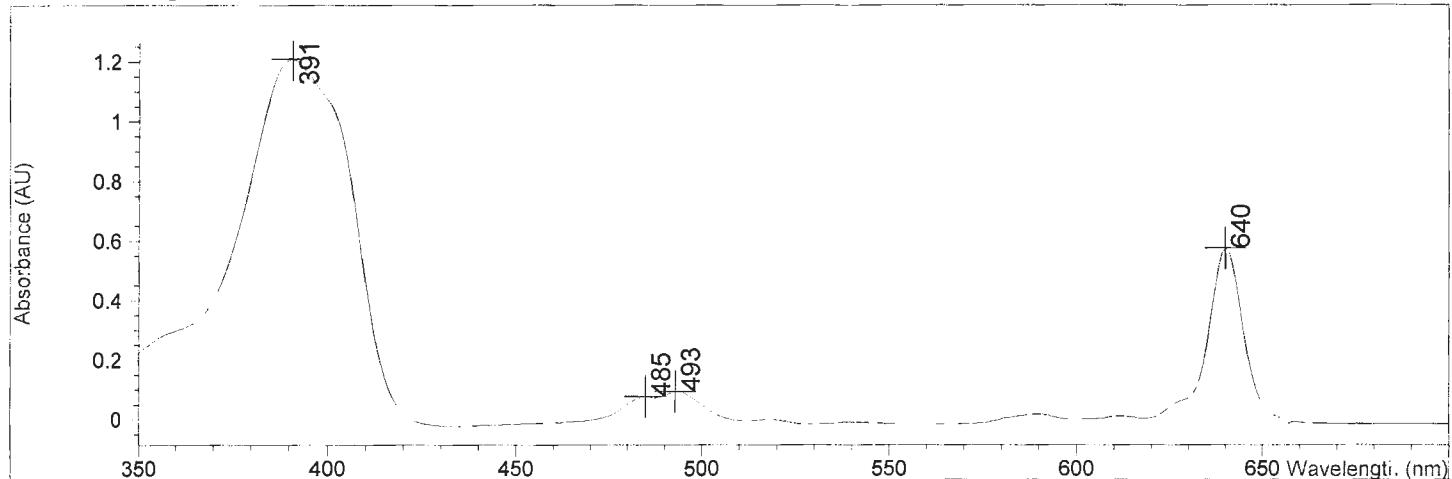
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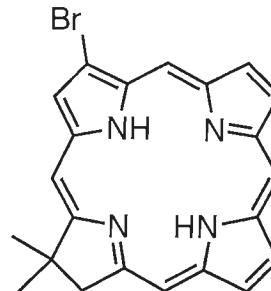


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