

**Supporting Information for “NDI and DAN DNA: Nucleic Acid Directed Assembly of NDI
and DAN”**

*Brian A. Ikkanda, Stevan A. Samuel, and Brent L. Iverson**

Department of Chemistry, The University of Texas at Austin, Austin, Texas, 78712

E-mail: iversonb@austin.utexas.edu

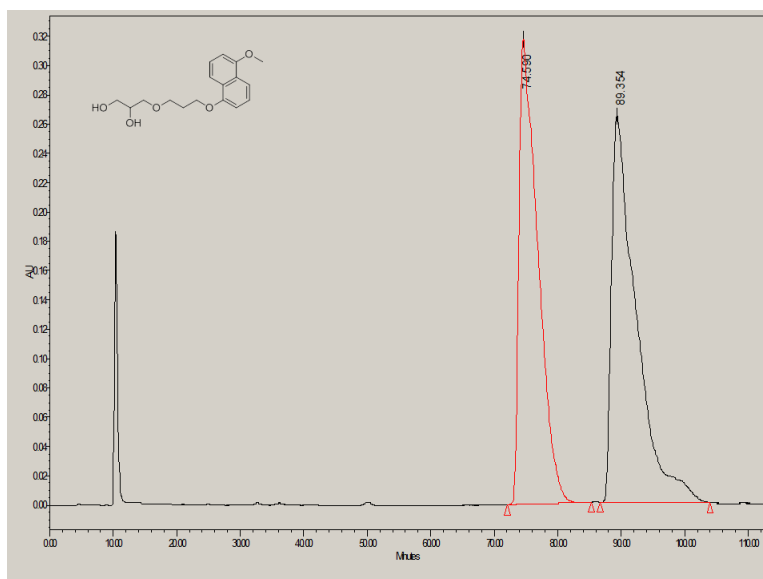
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General Procedures and Reagents

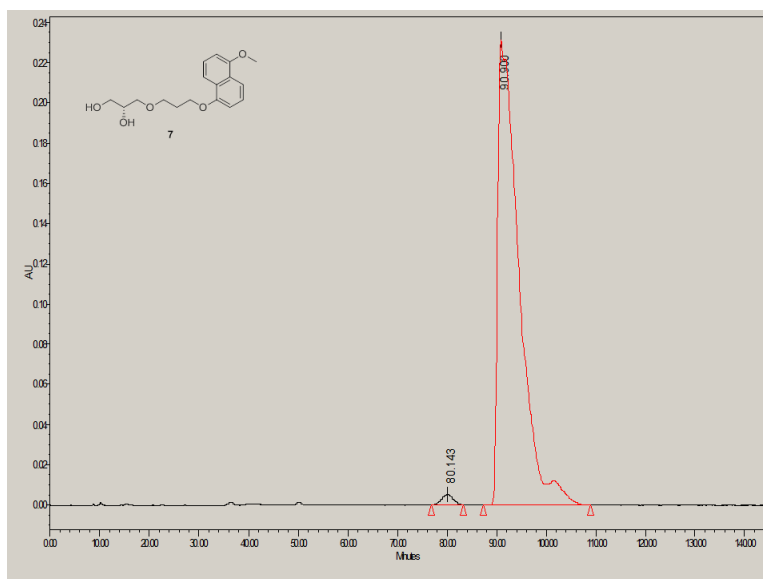
High-resolution mass spectra were analyzed by Q-TOF. Reactions were performed under an atmosphere of argon unless otherwise specified. All microwave reactions were conducted in a C.E.M. MARS microwave reactor at a power of 300W in an open vessel or C.E.M. Explorer 48 microwave reactor in a sealed vessel at a power of 300W.

Figure S1. ^a Chiral HPLC Chromatograms of 7 and 10.

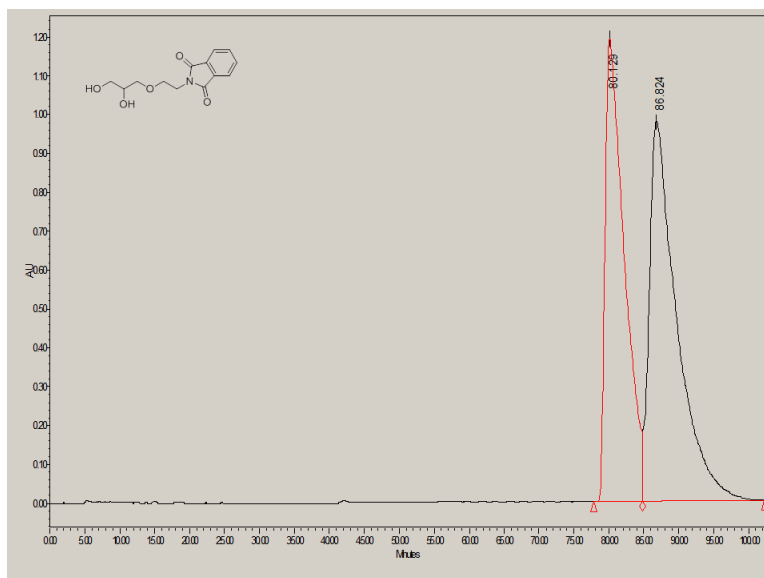


Racemic Sample

	Retention Time	Area	% Area	Height
1	80.129	226355648	45.90	1189756
2	86.824	266826823	54.10	976579

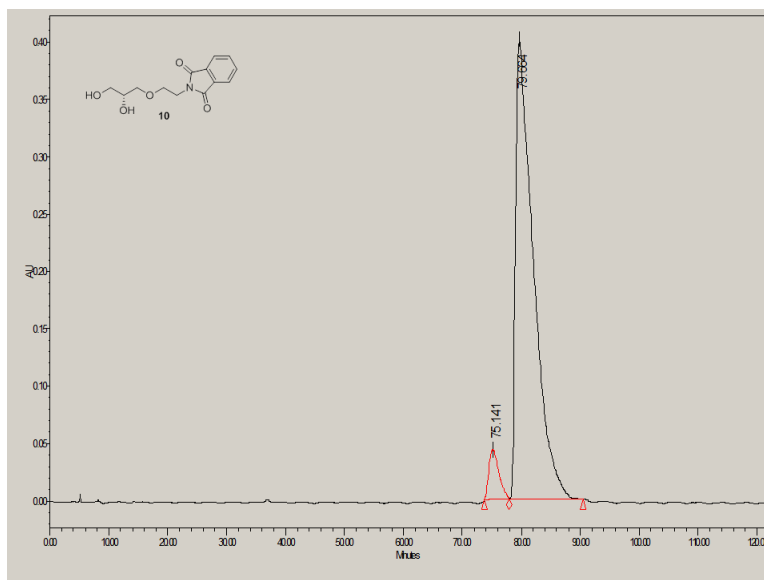


	Retention Time	Area	% Area	Height
1	75.141	5187344	5.82	43704
2	79.664	83915537	94.18	401161



Racemic Sample

	Retention Time	Area	% Area	Height
1	74.590	65668561	48.10	316870
2	89.354	70865309	51.90	263866



	Retention Time	Area	% Area	Height
1	80.143	824716	1.19	5290
2	90.900	68763657	98.81	231161

^a Enantiomeric excess was determined by HPLC on a Chiralcel ODH column (0.46 cm I.D. x 25 cmL); eluent, hexane:*i*-PrOH, 95:5 v/v; flow rate, 1.0 mL/min; UV at 254 nm; room temperature.

Figure S2. ^1H NMR (CDCl_3) of **1**

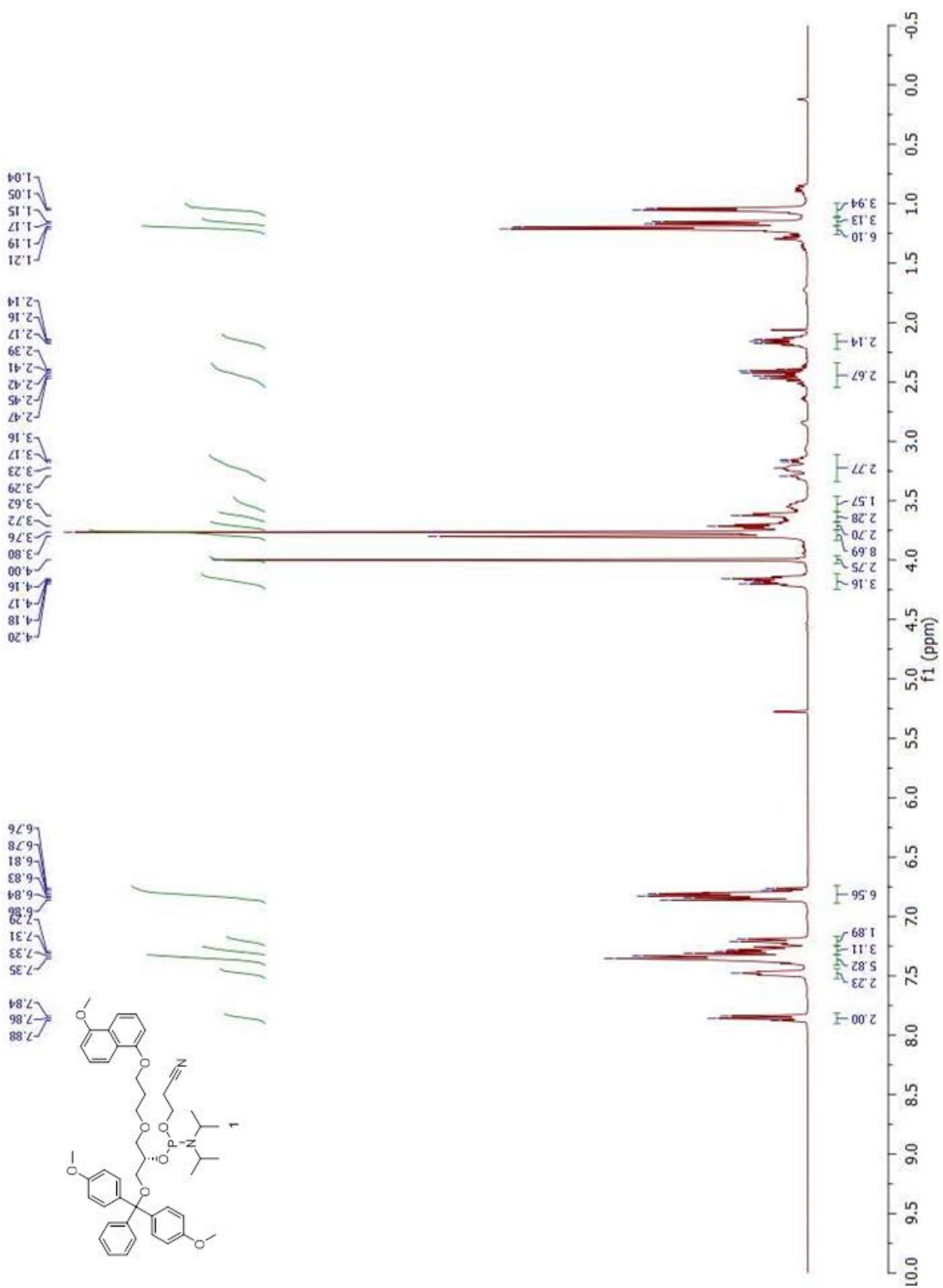


Figure S3. ^1H NMR (CDCl_3) of **2**

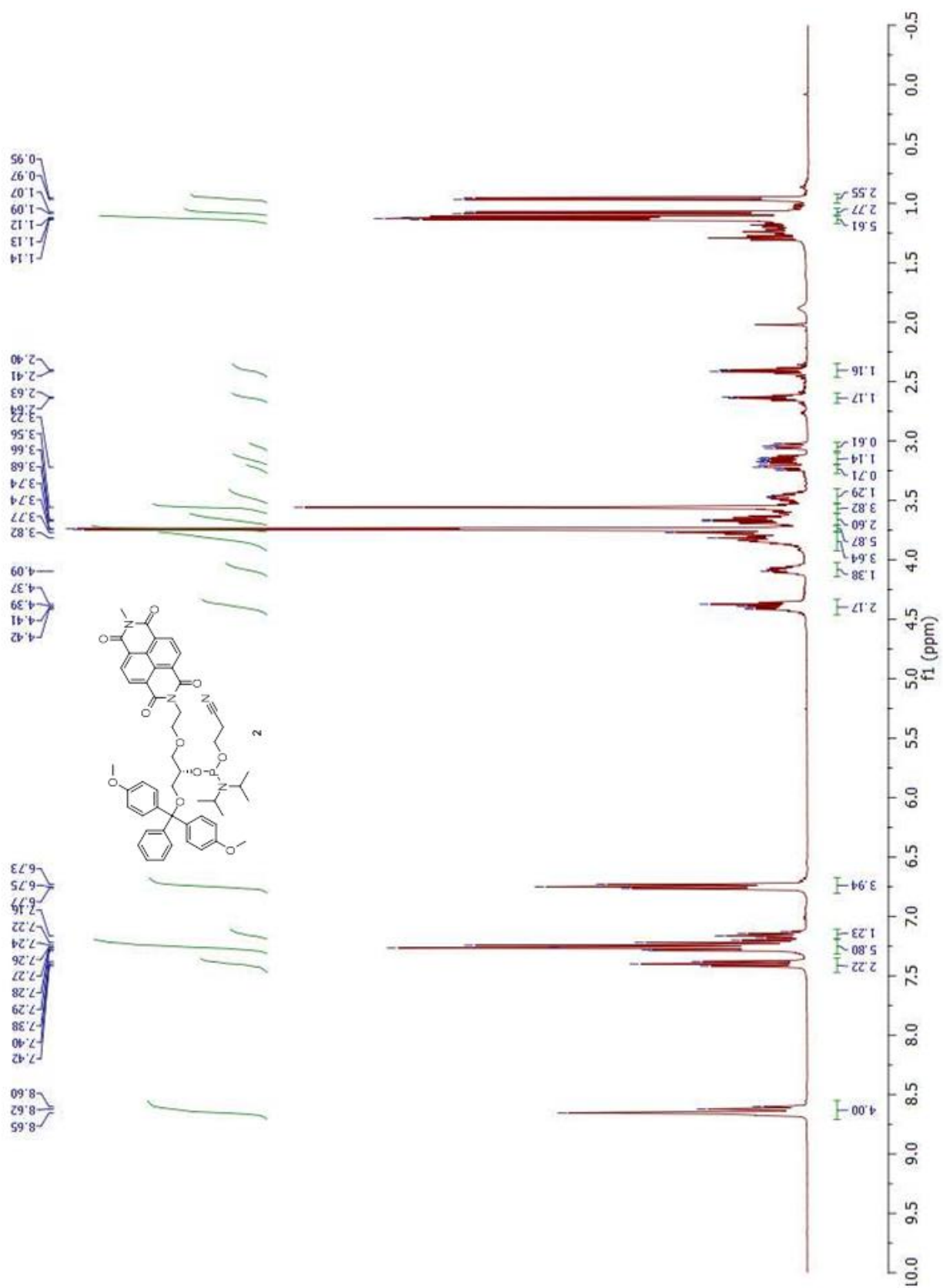


Figure S4. ^1H NMR (CDCl_3) of **3**

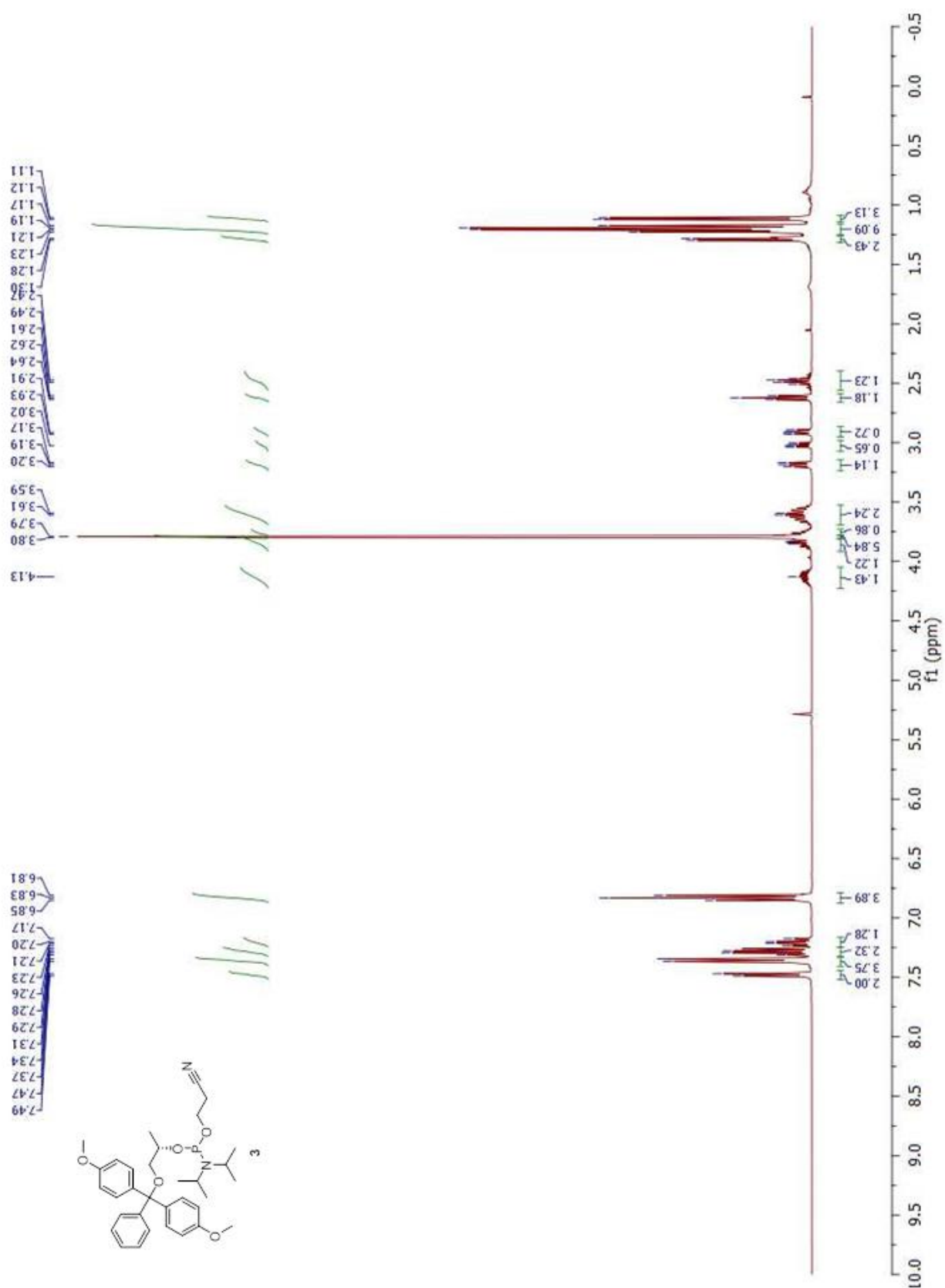


Figure S5. ^1H NMR (CDCl_3) of **5**

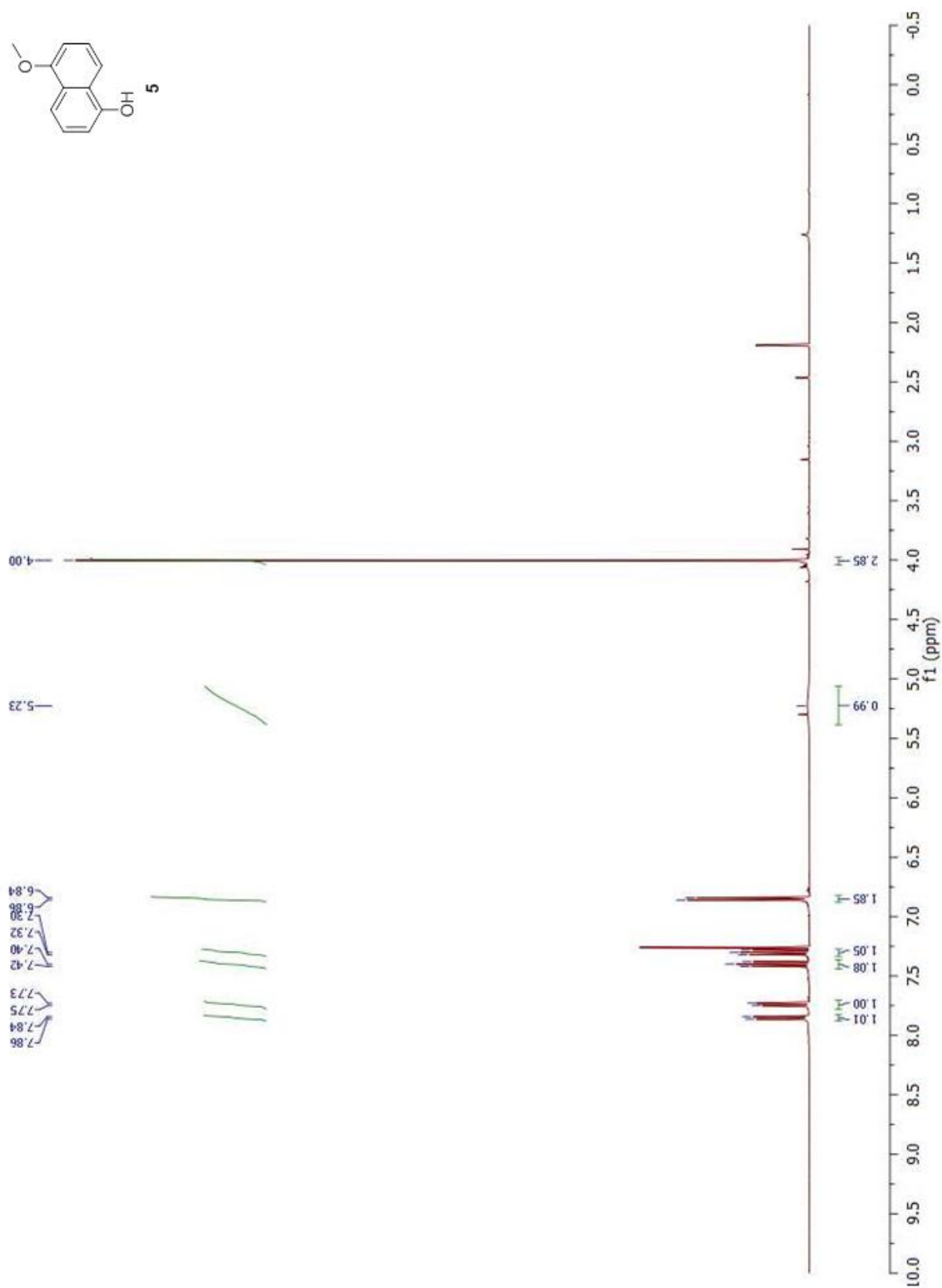


Figure S6. ^1H NMR (CDCl_3) of **6**

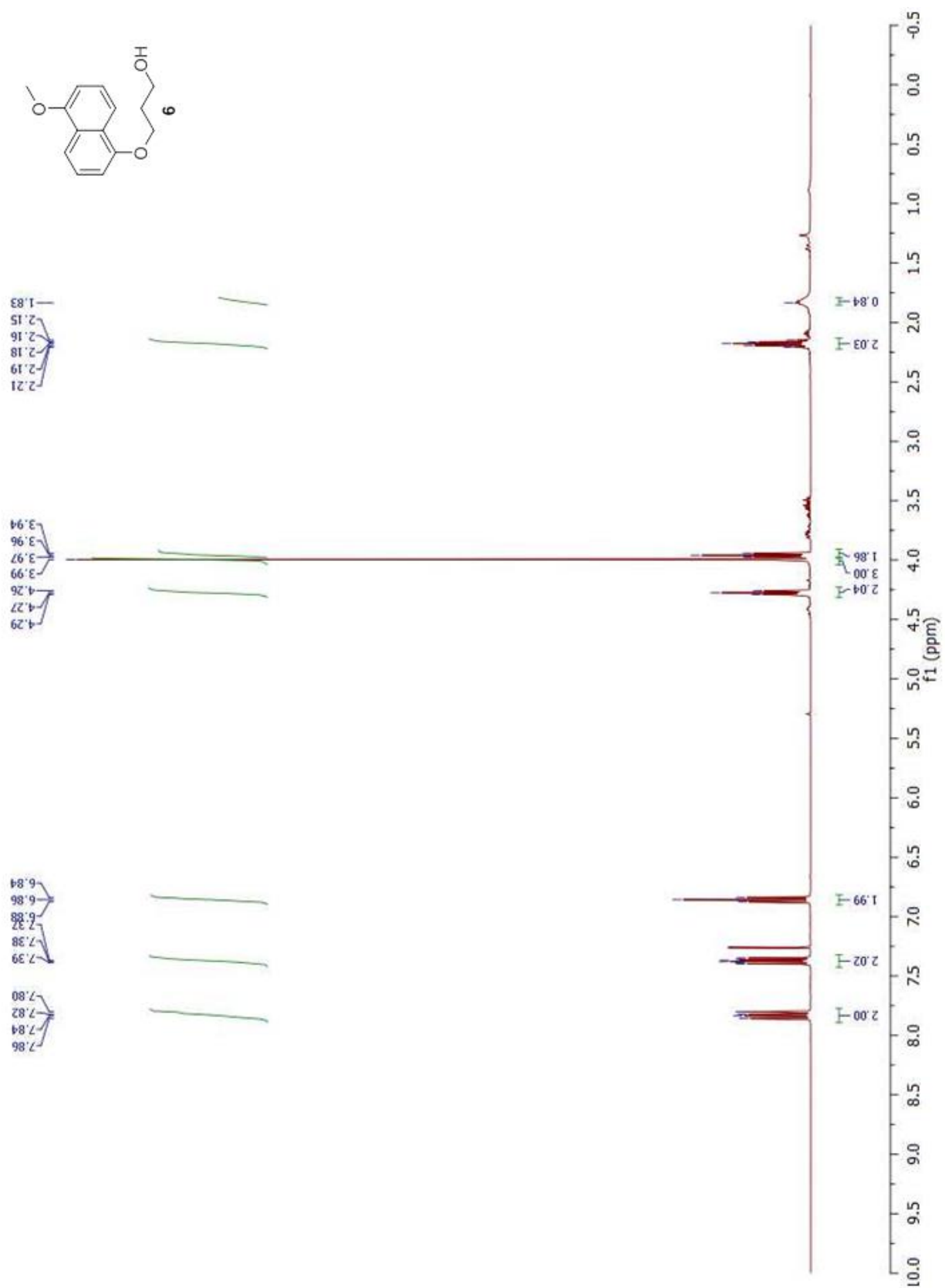


Figure S7. ^1H NMR (CDCl_3) of **7**

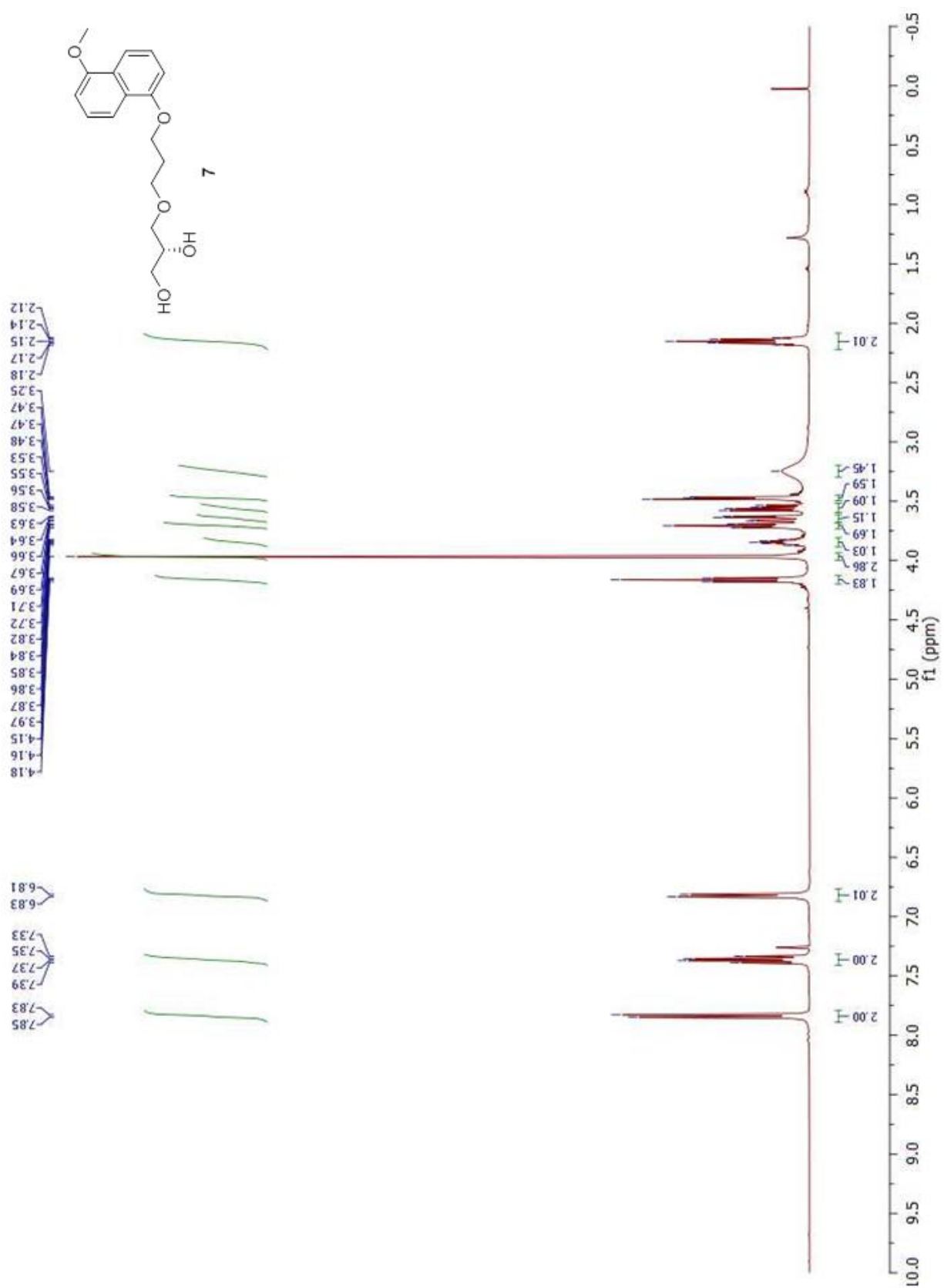


Figure S8. ^1H NMR (CDCl_3) of **8**

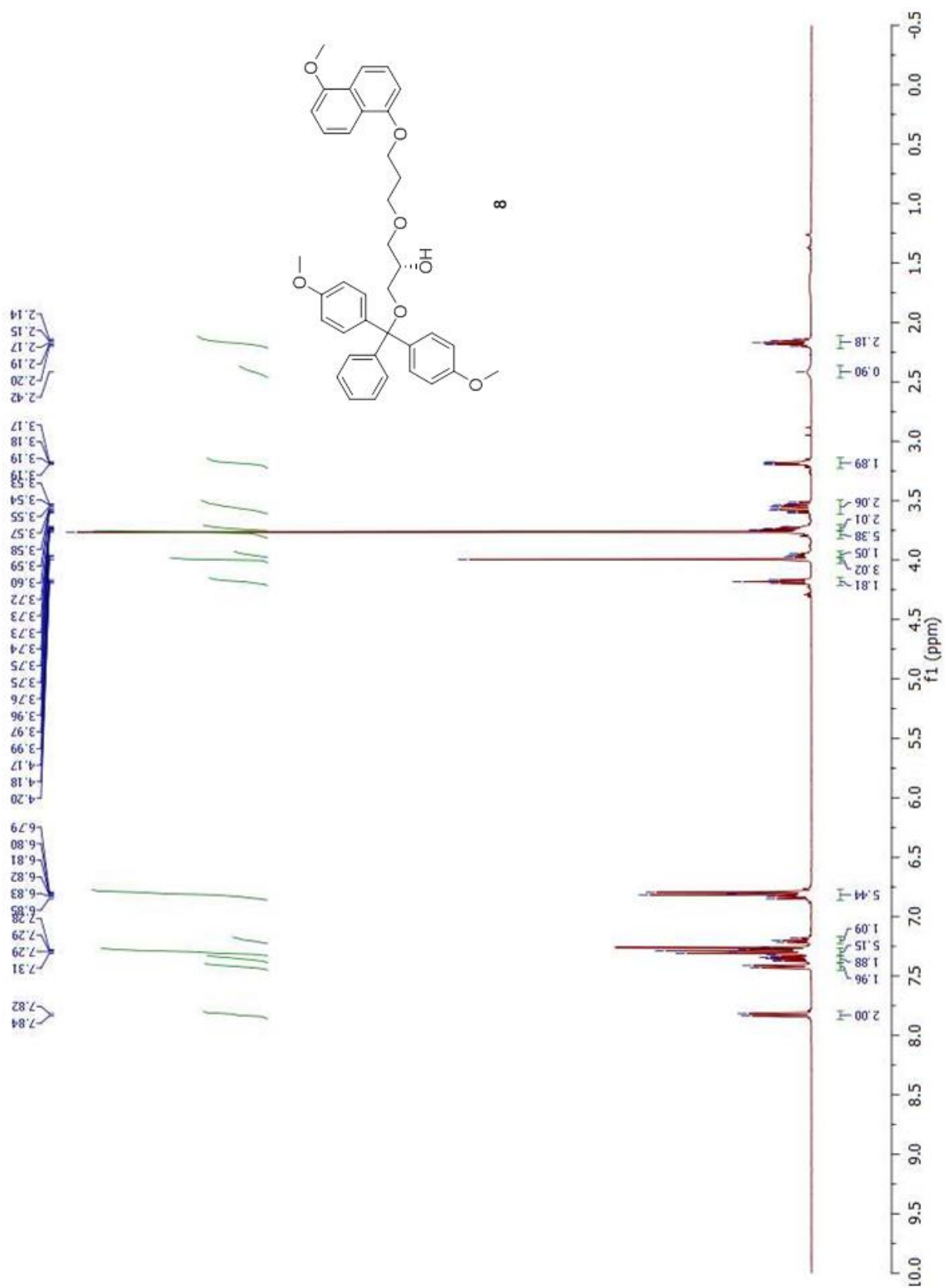
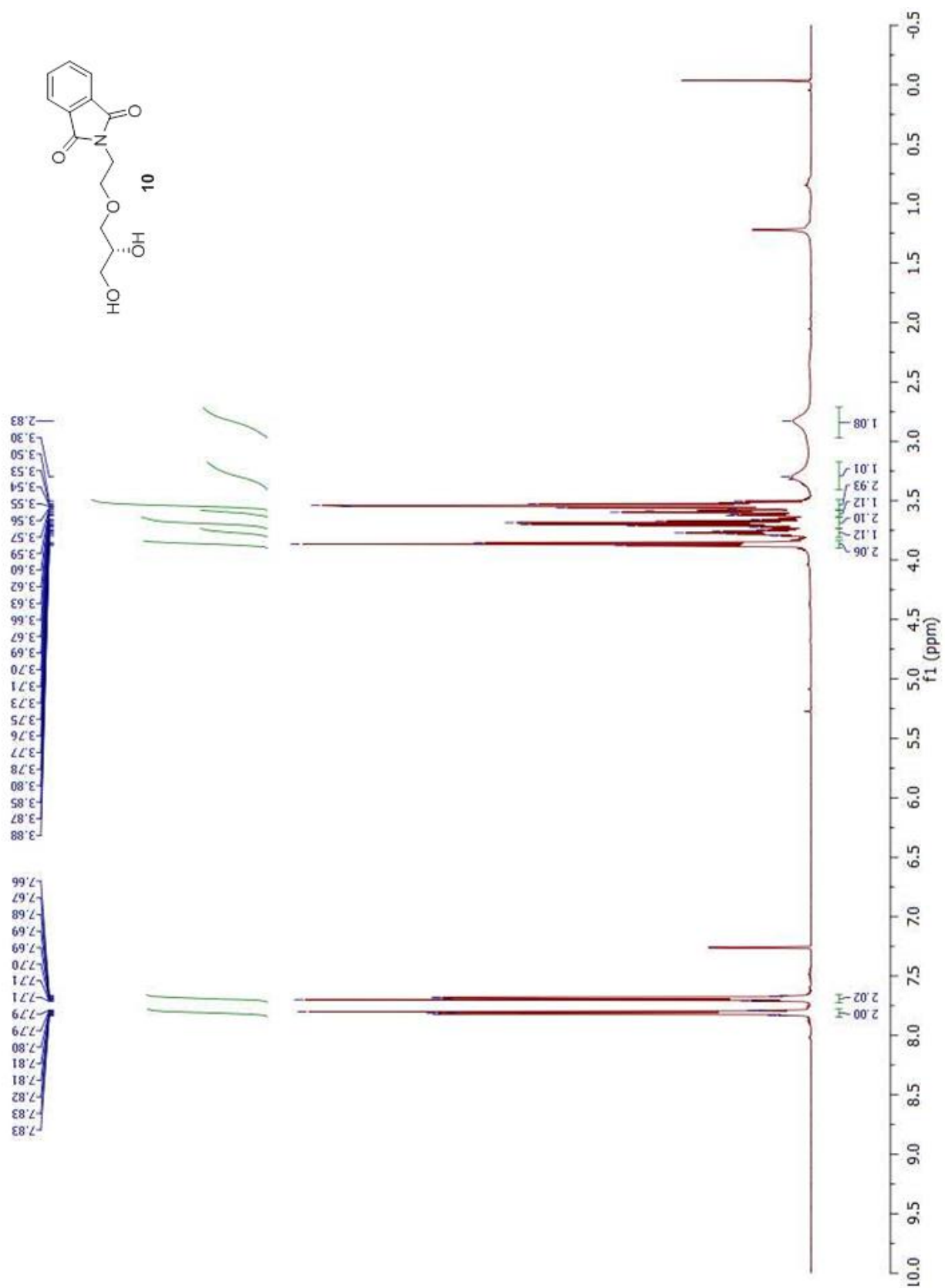


Figure S9. ^1H NMR (CDCl_3) of **10**



Chemical structure of compound 11 is shown in the top left. The ¹H NMR spectrum (CDCl₃) is displayed below, with chemical shifts (ppm) on the x-axis ranging from 0.5 to 10.0. Integration values are provided for several peak regions.

Chemical structure of 11: COc1ccc(cc1)Oc2ccc(cc2)C(c3ccccc3)(c4cc(OC)ccc4)COC[C@H](O)COc5cc6c(cc5)nc(=O)c7ccccc76=O

¹H NMR spectrum (CDCl₃) data:

Chemical Shift (ppm)	Integration
0.61	0.61
2.24	2.04
2.56	1.04
3.61	2.04
3.73	3.73
5.72	2.02
5.77	0.99
7.21	2.00

Figure S11. ^1H NMR (CDCl_3) of **12**

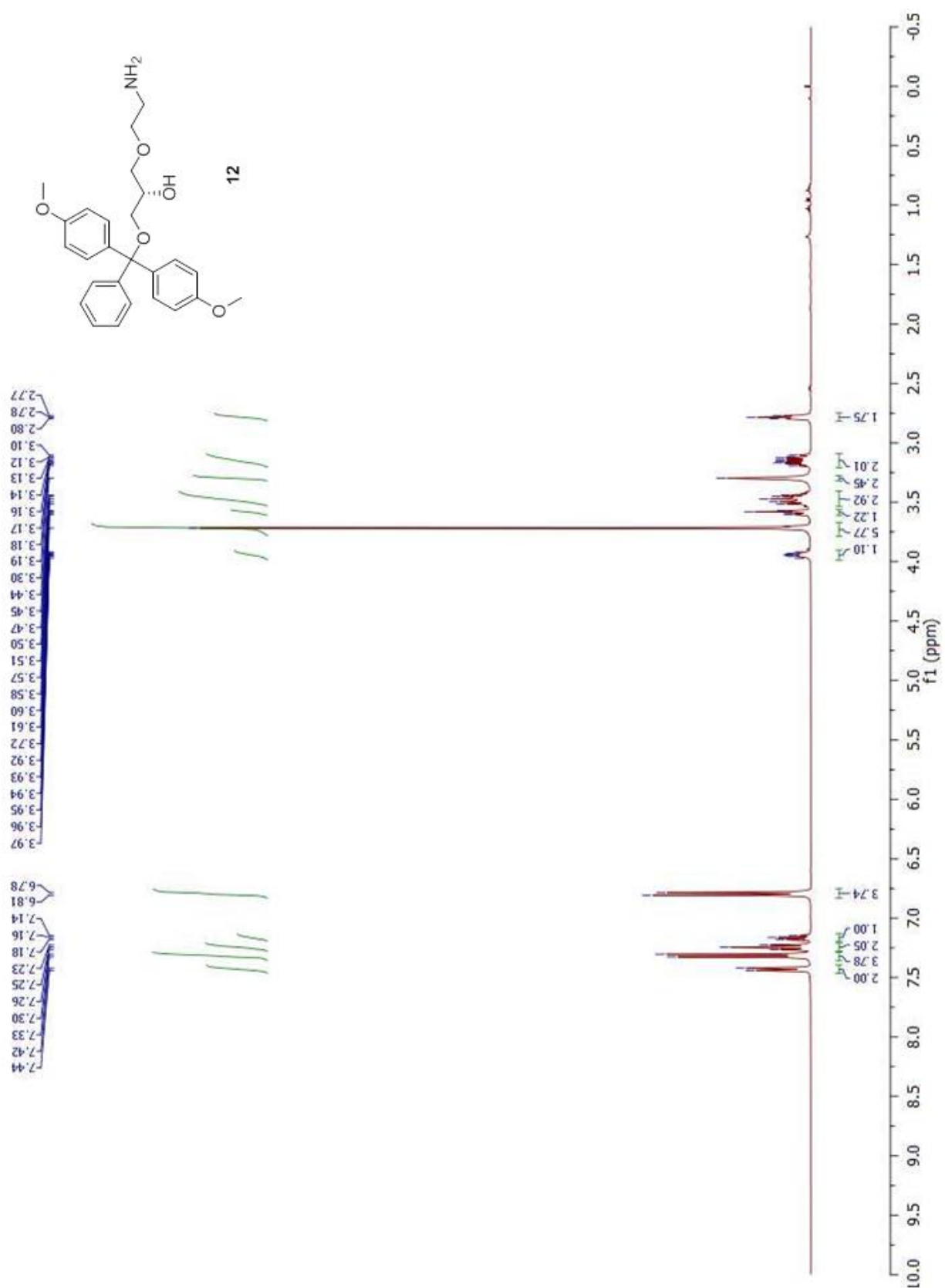


Figure S12. ^1H NMR (CDCl_3) of **14**

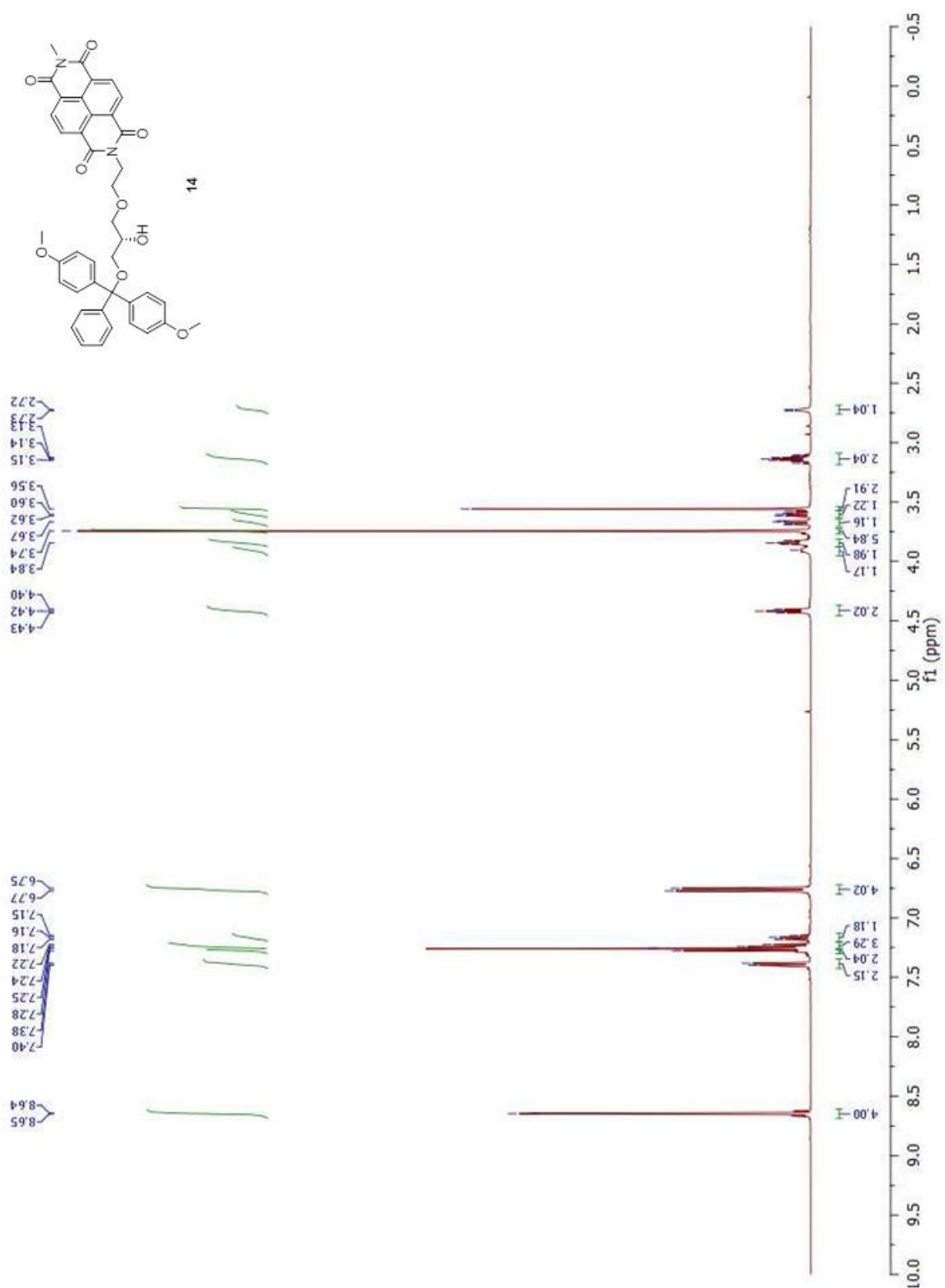


Figure S13. ^{13}C NMR (CDCl_3) of **1**

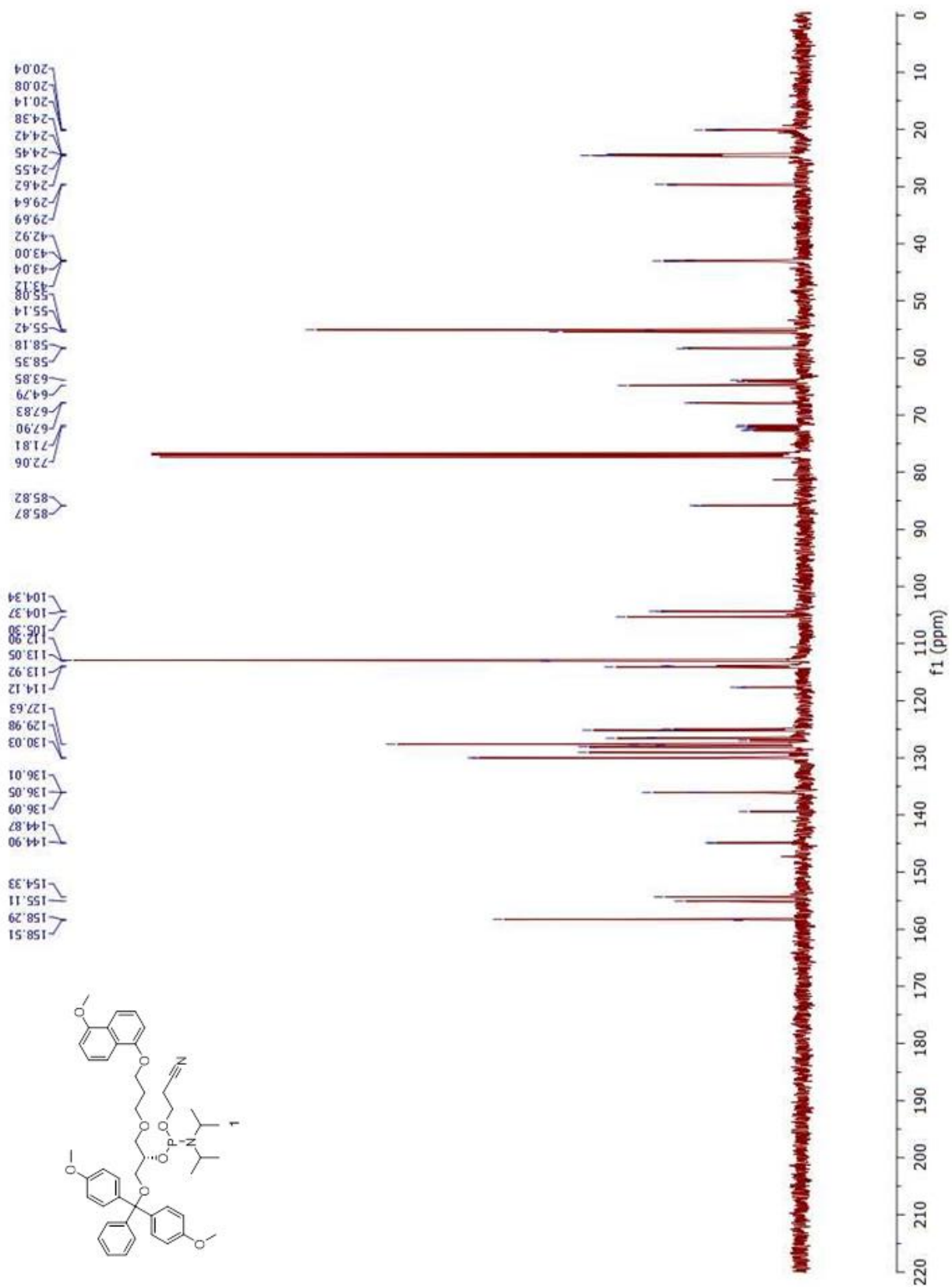


Figure S14. ^{13}C NMR (CDCl_3) of **2**

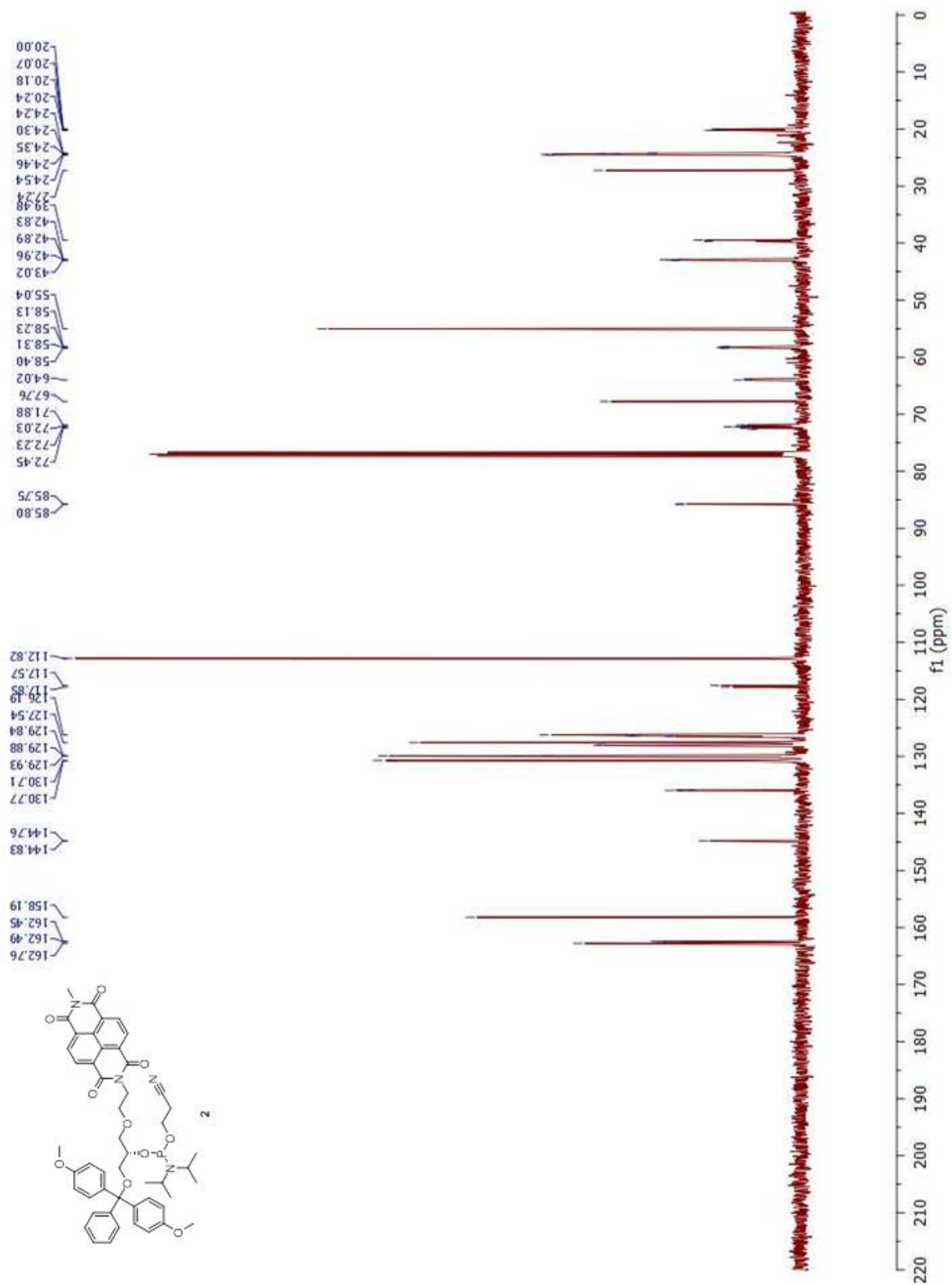


Figure S15. ^{13}C NMR (CDCl_3) of **3**

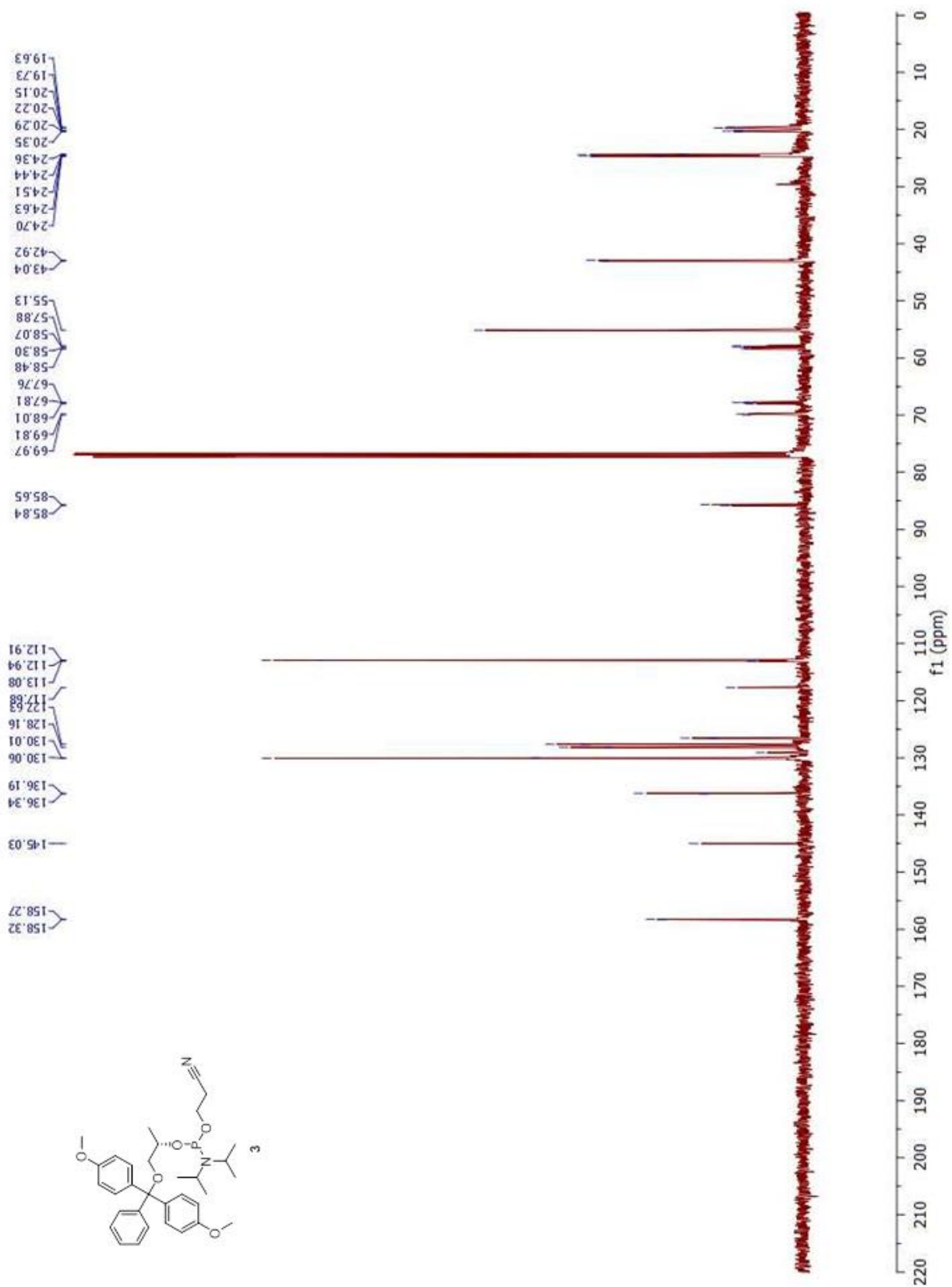


Figure S16. ^{13}C NMR (CDCl_3) of **5**

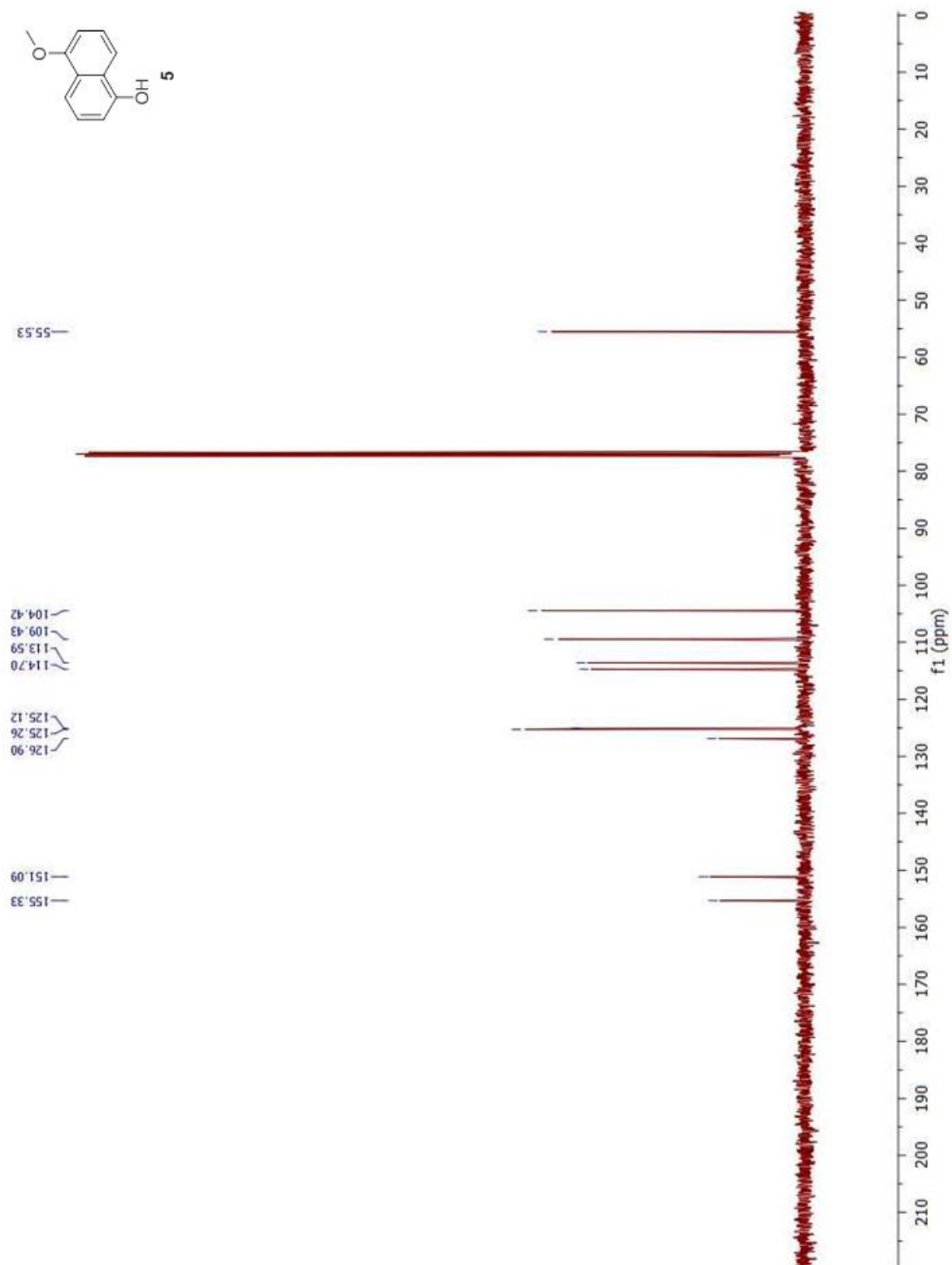


Figure S17. ^{13}C NMR (CDCl_3) of **6**

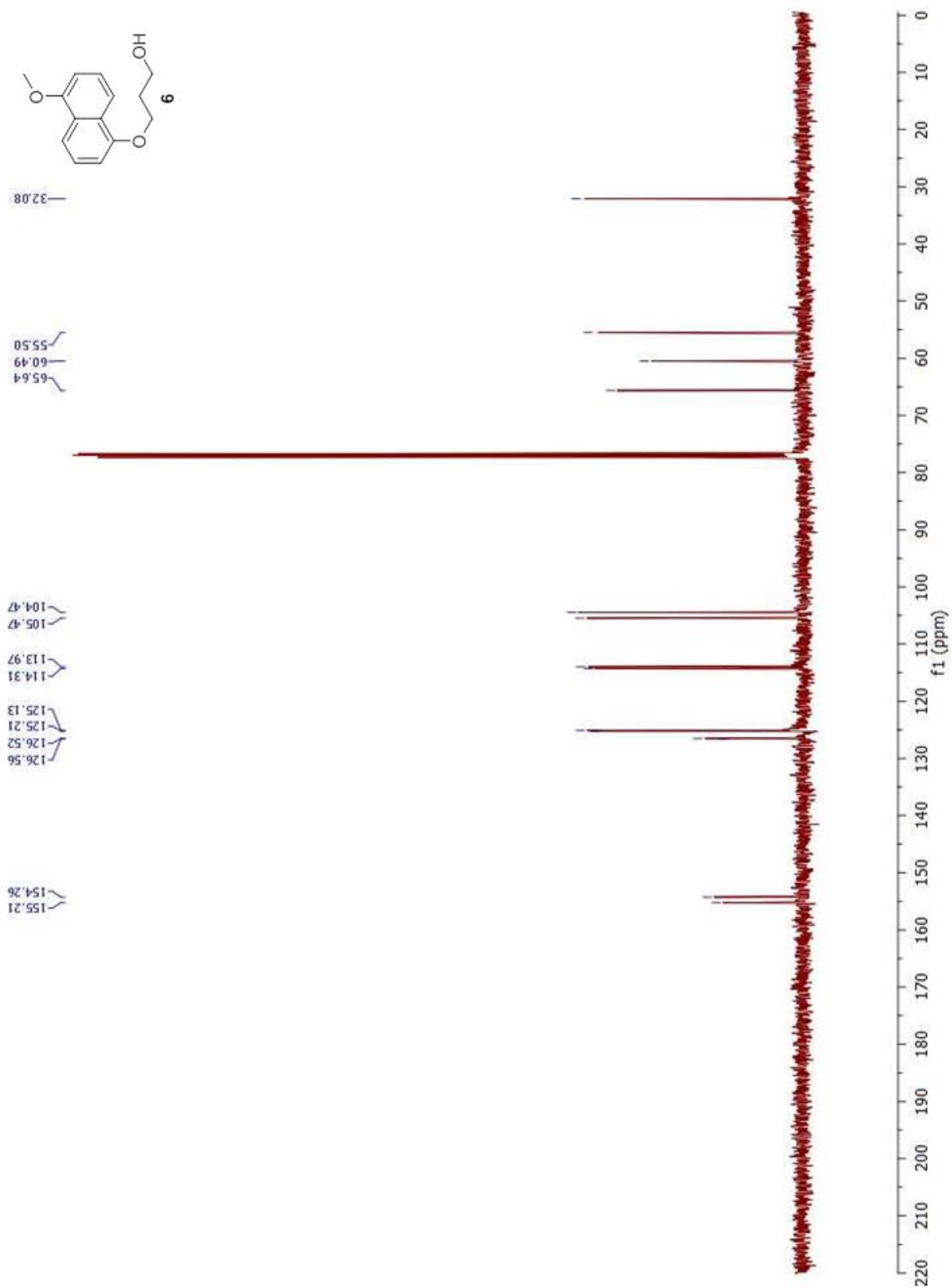


Figure S18. ^{13}C NMR (CDCl_3) of **7**

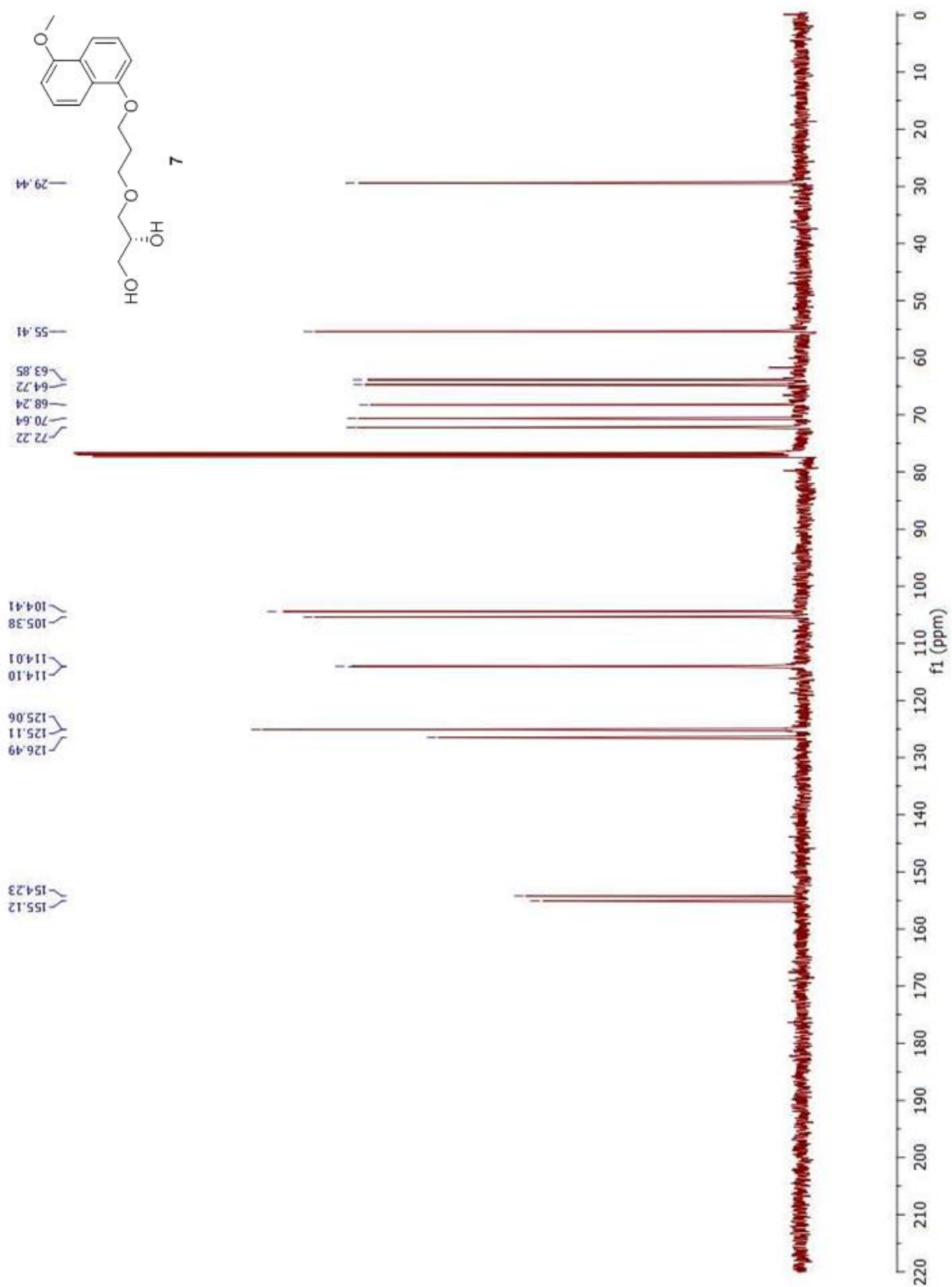


Figure S19. ^{13}C NMR (CDCl_3) of **8**

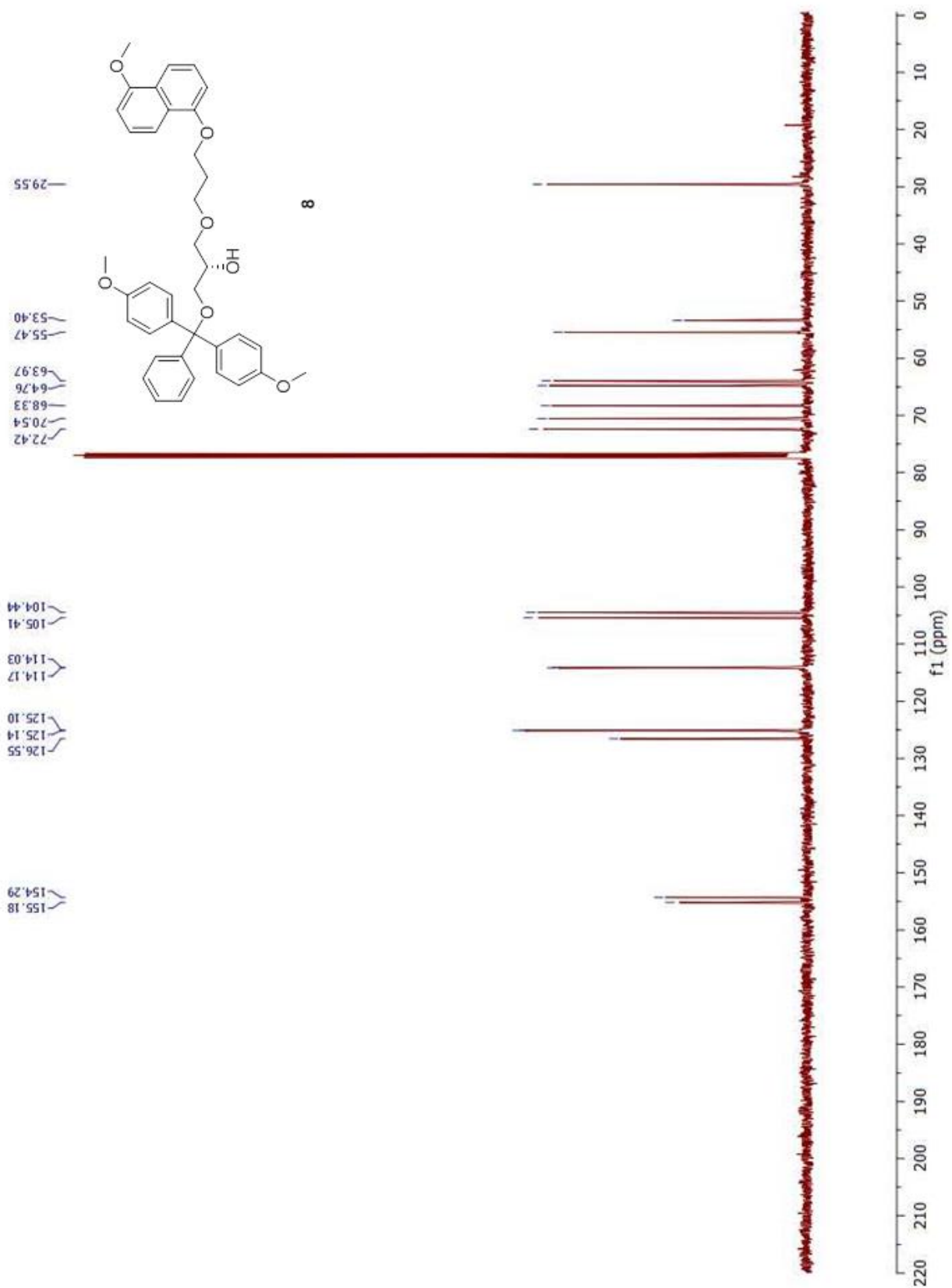


Figure S20. ^{13}C NMR (CDCl_3) of **10**

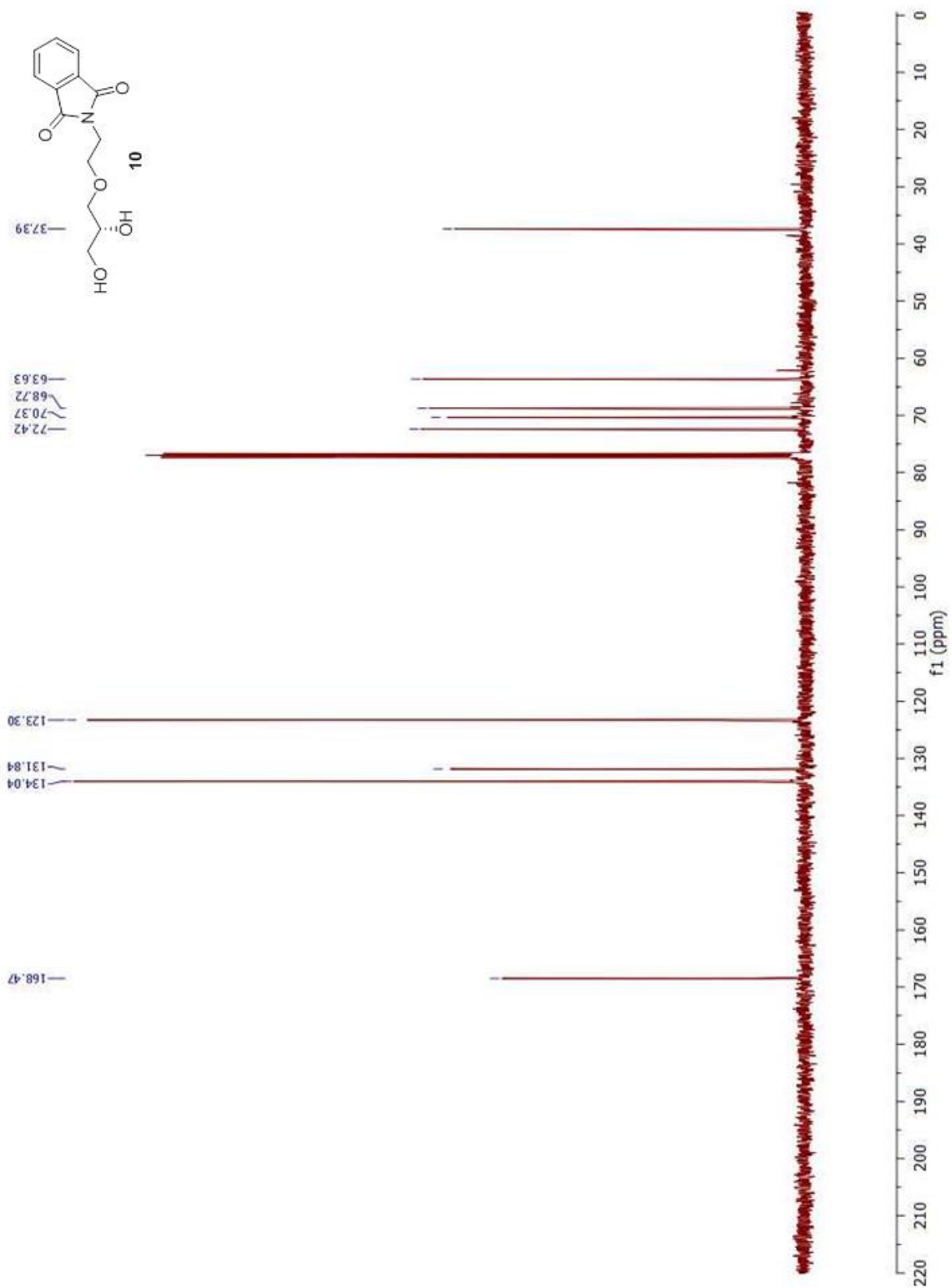


Figure S21. ^{13}C NMR (CDCl_3) of **11**

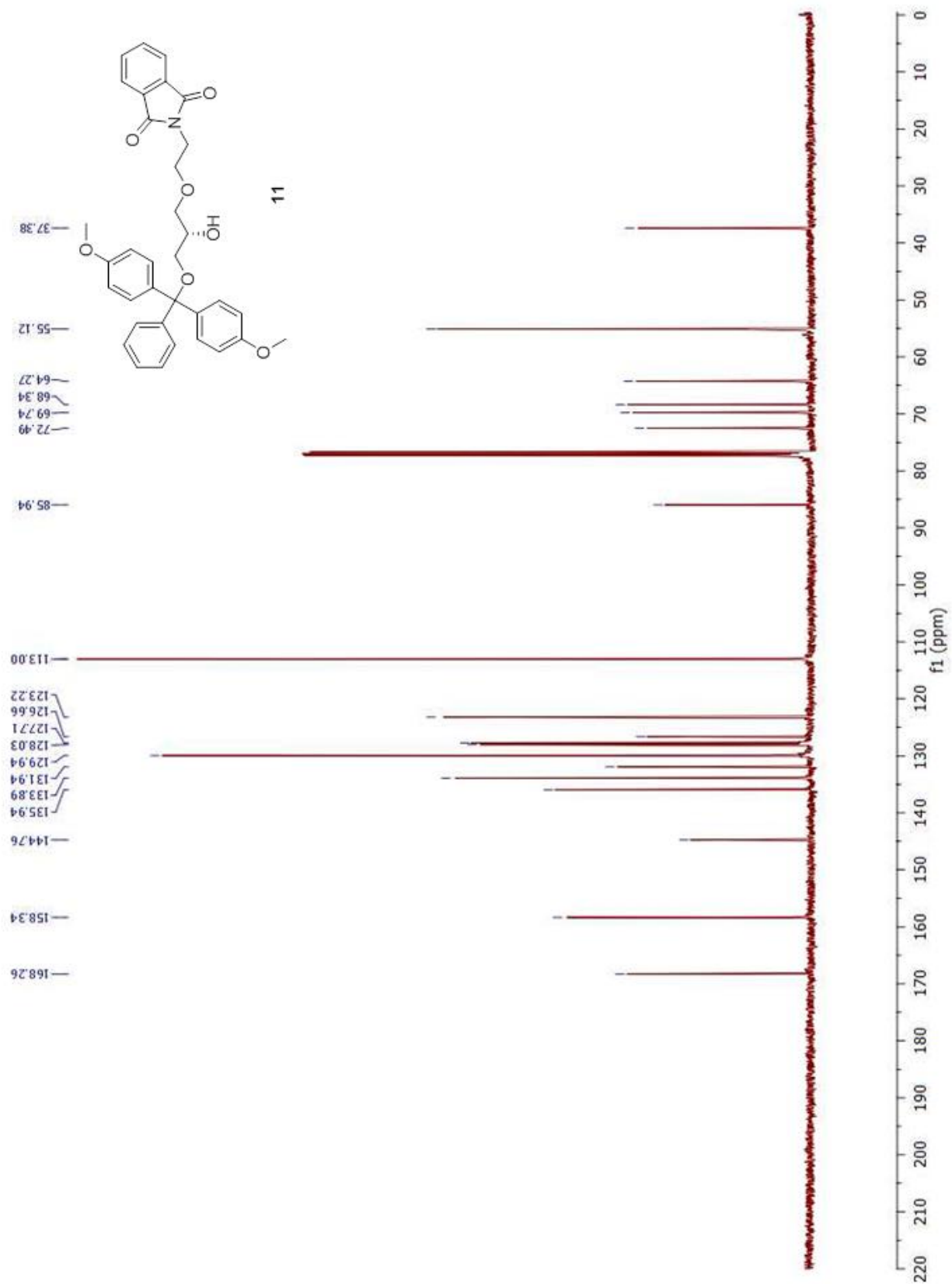


Figure S22. ^{13}C NMR (CDCl_3) of **12**

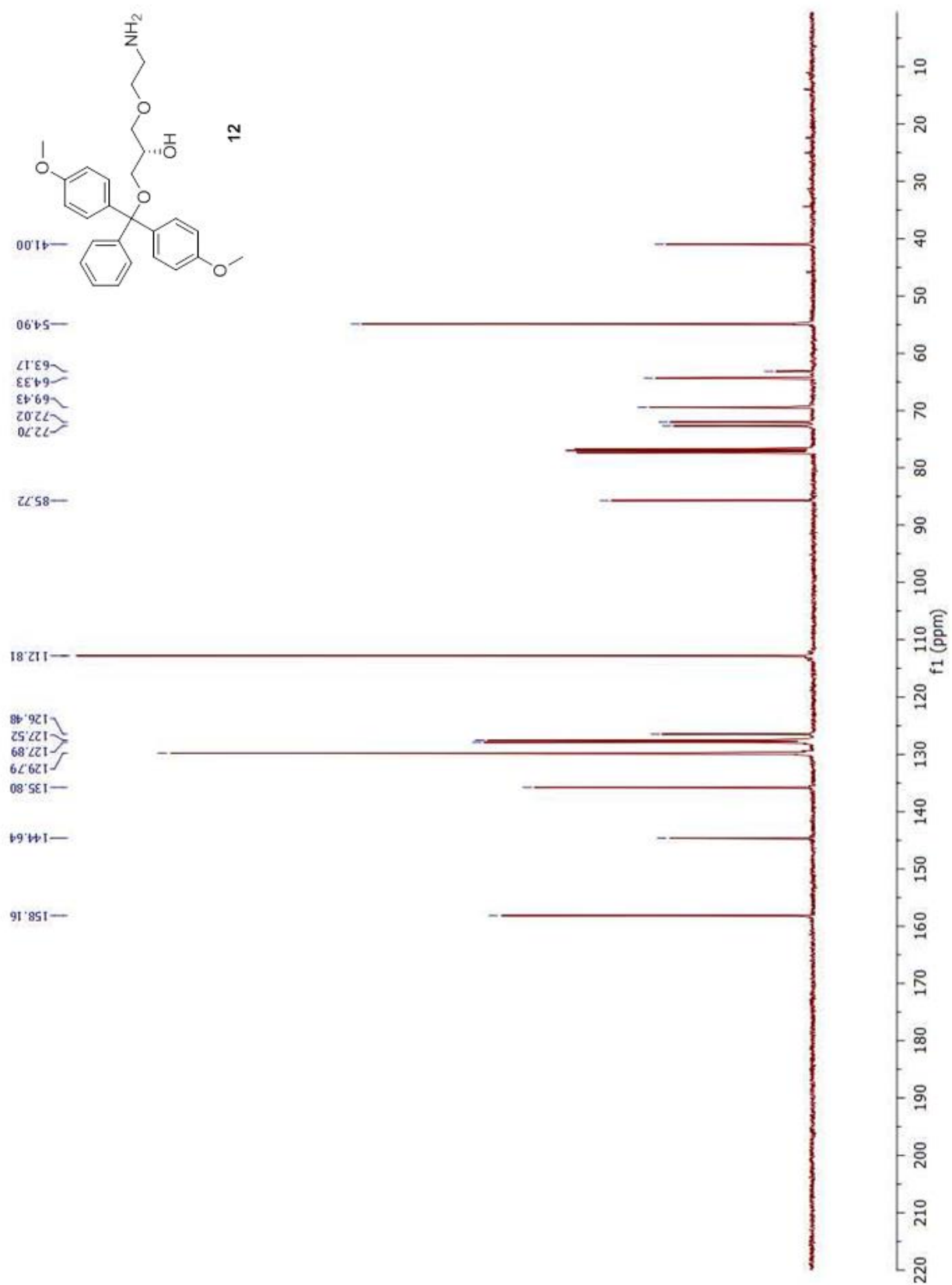


Figure S23. ^{13}C NMR (CDCl_3) of **14**

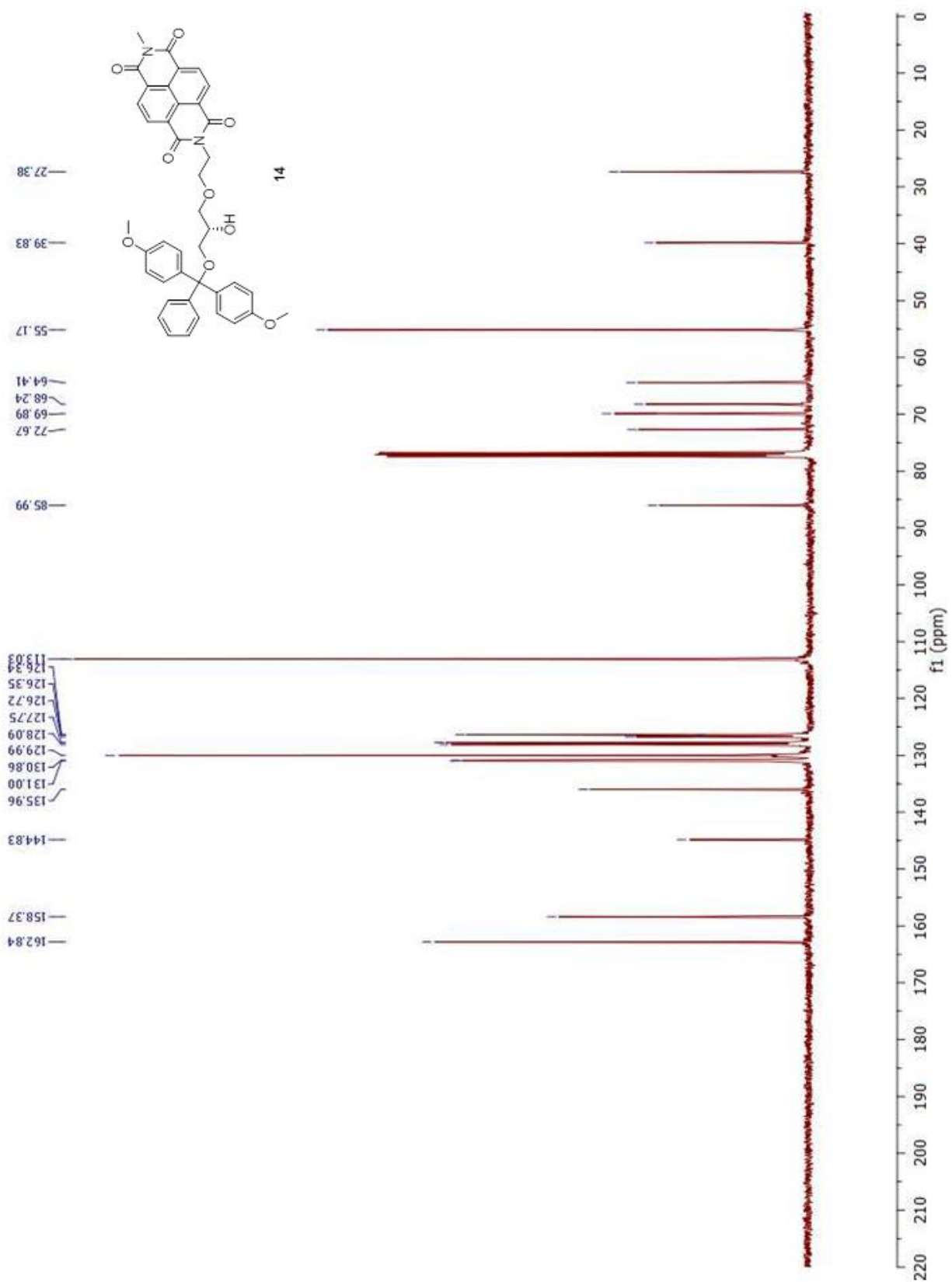


Figure S24. ^{31}P NMR (CDCl_3) of **1**

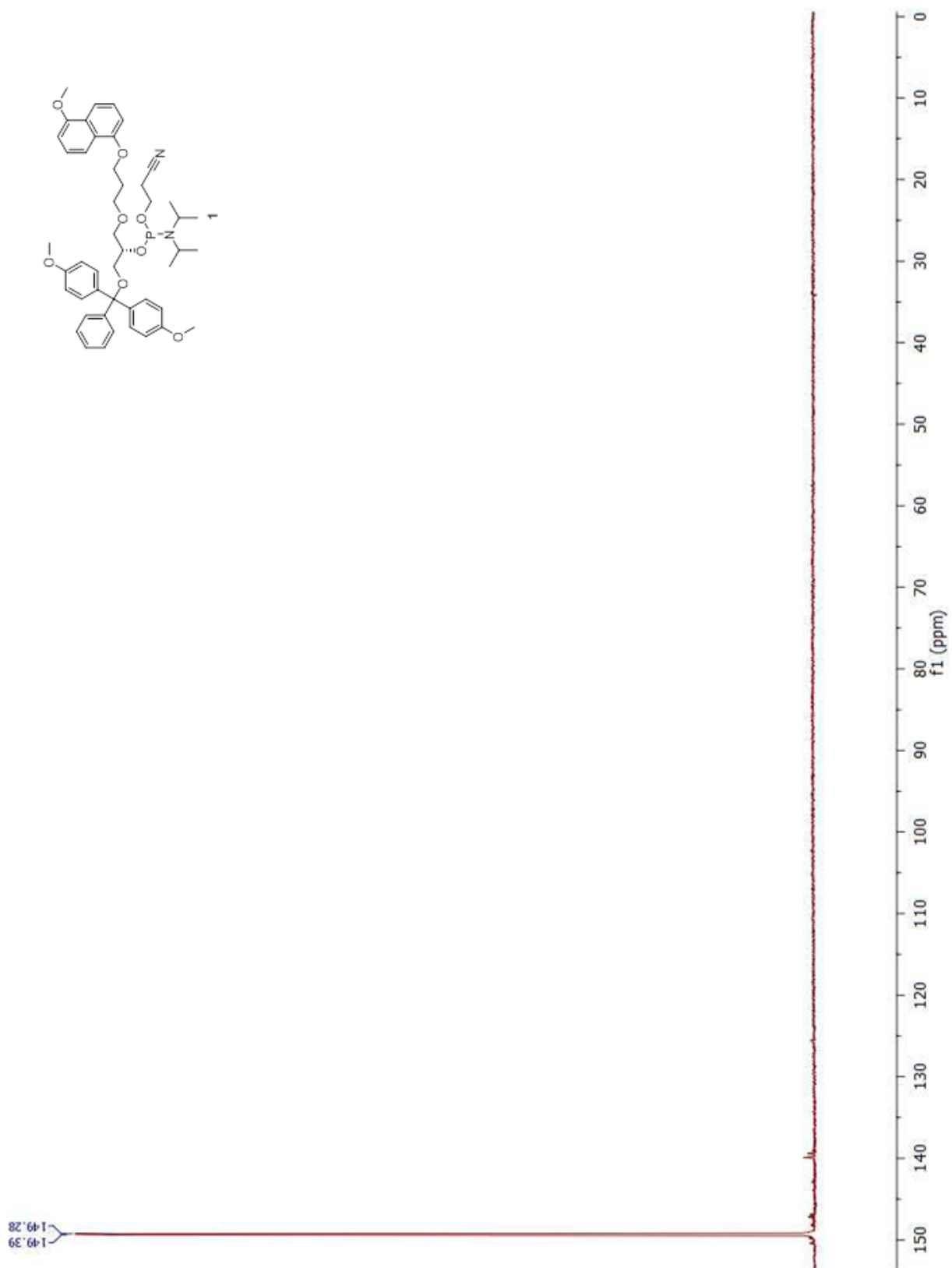
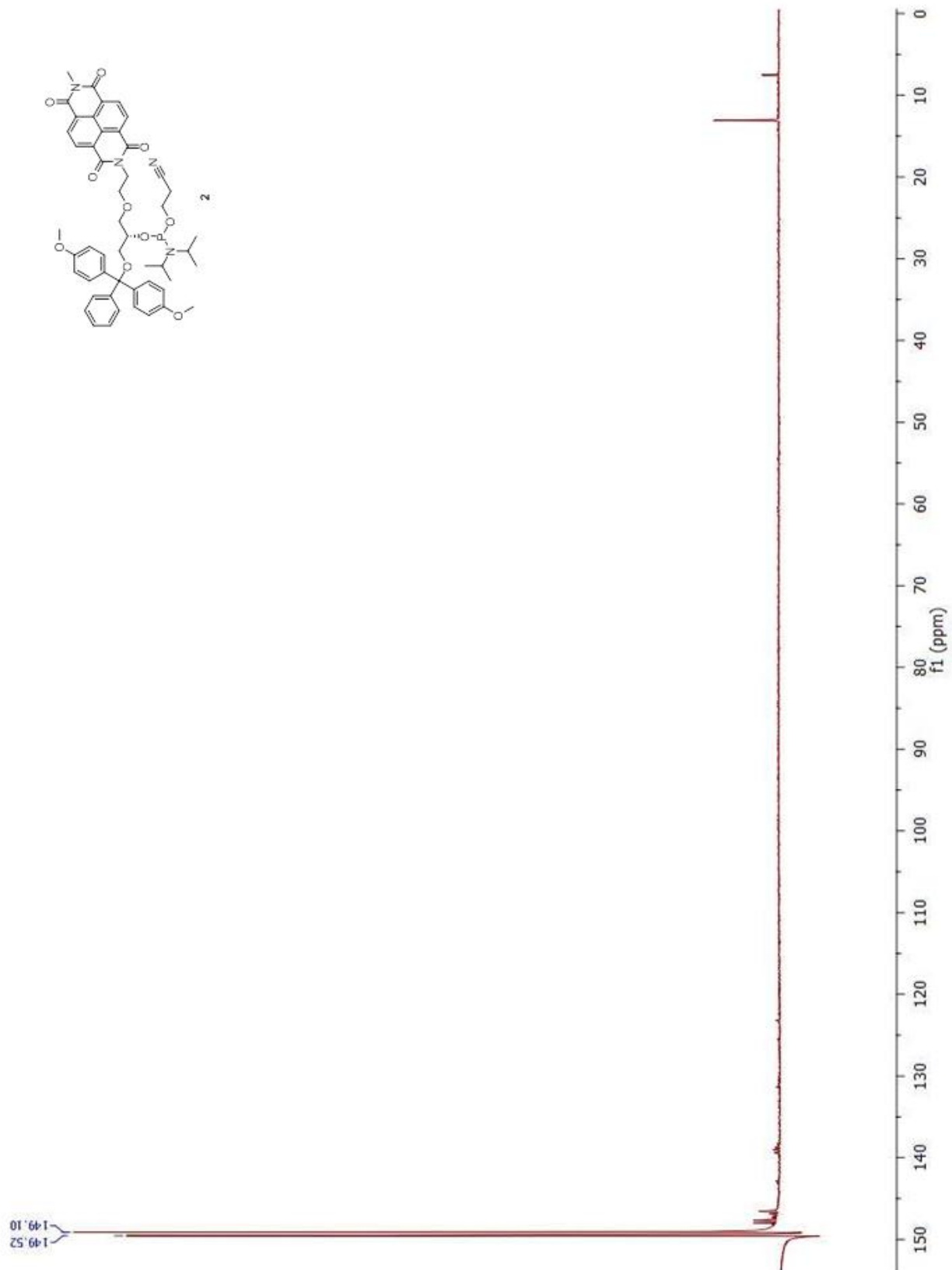


Figure S25. ^{31}P NMR (CDCl_3) of **2**



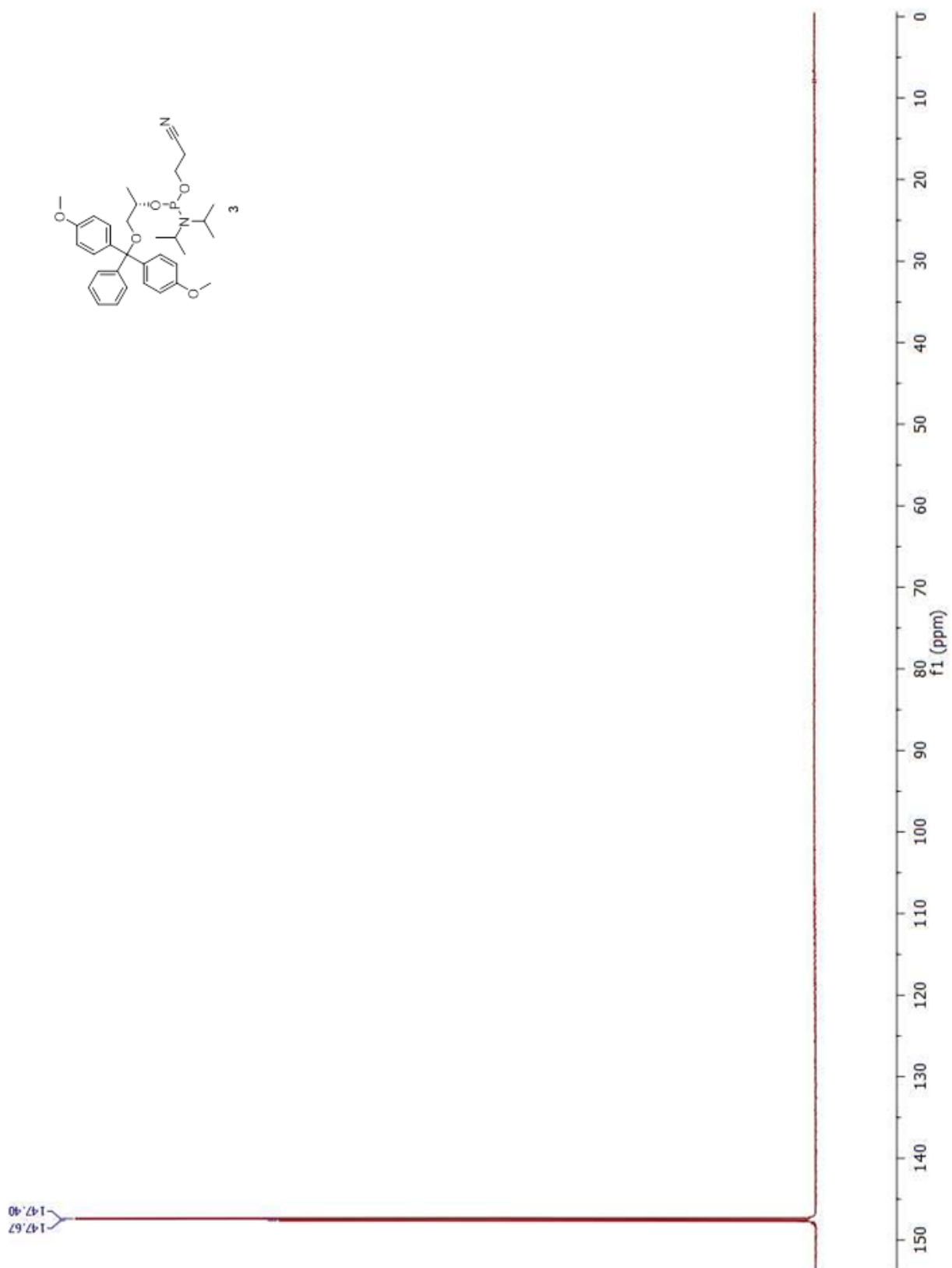
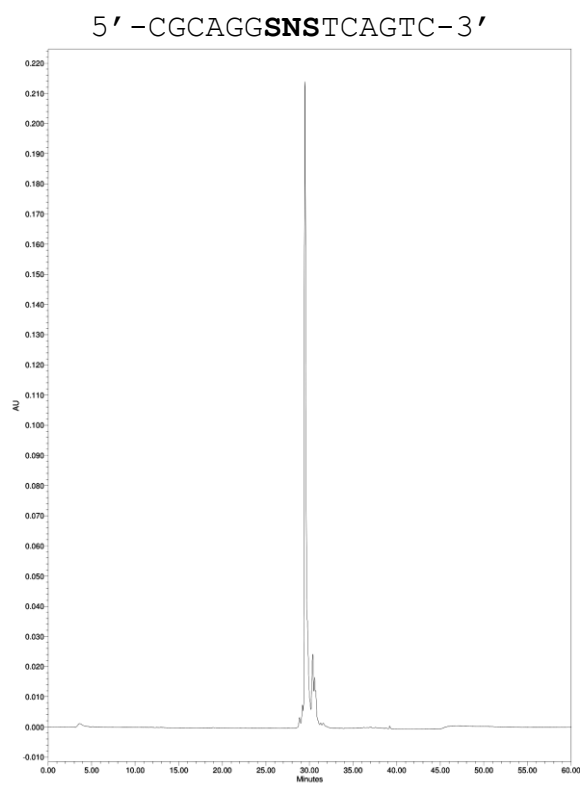
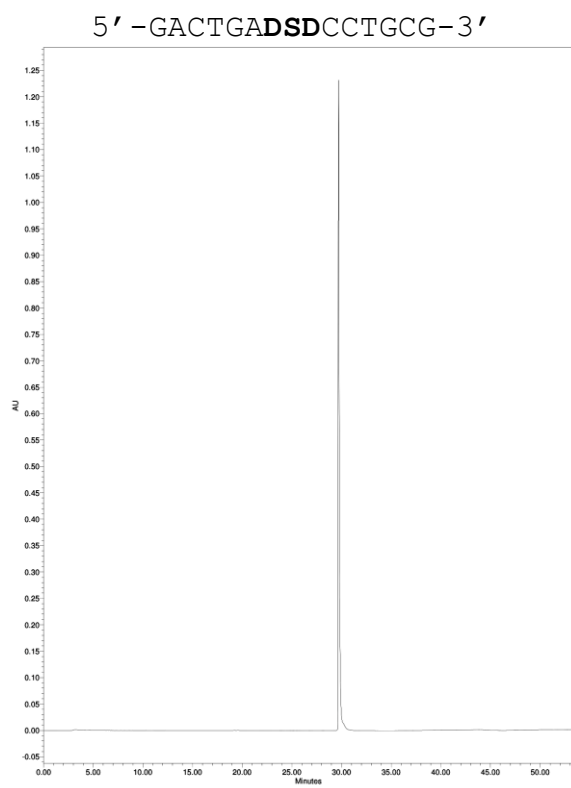
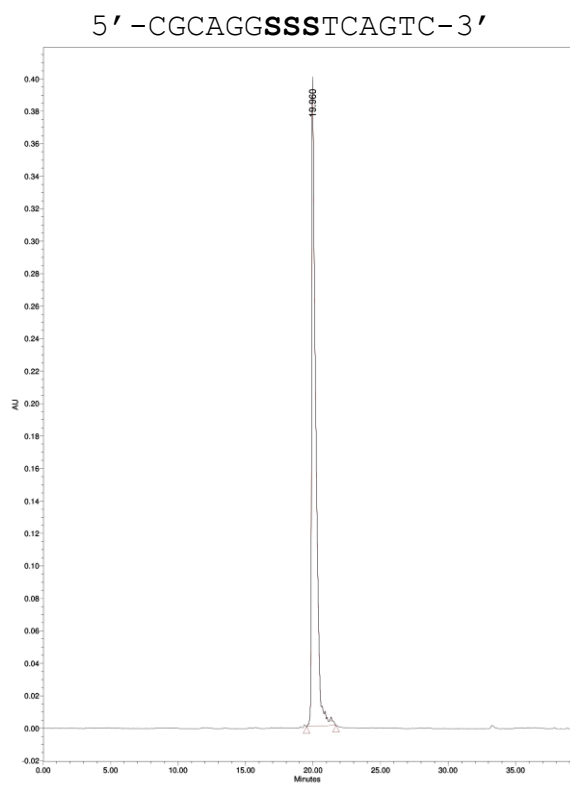
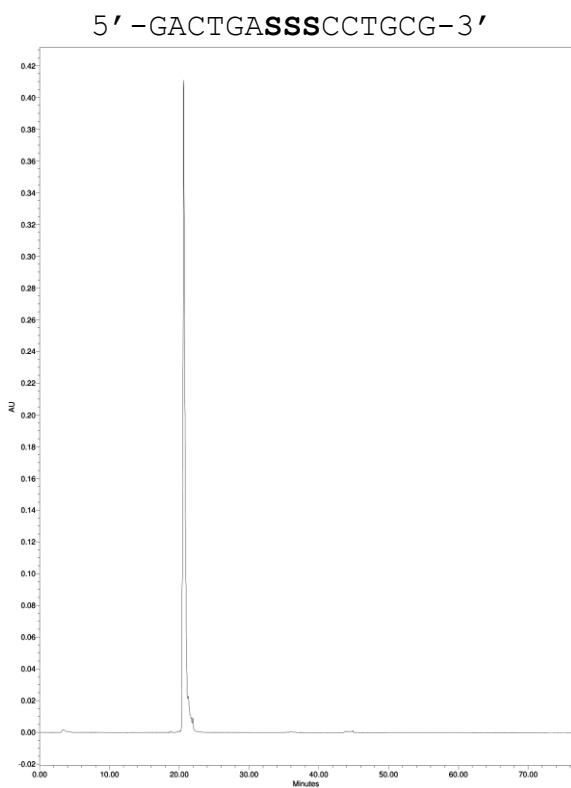
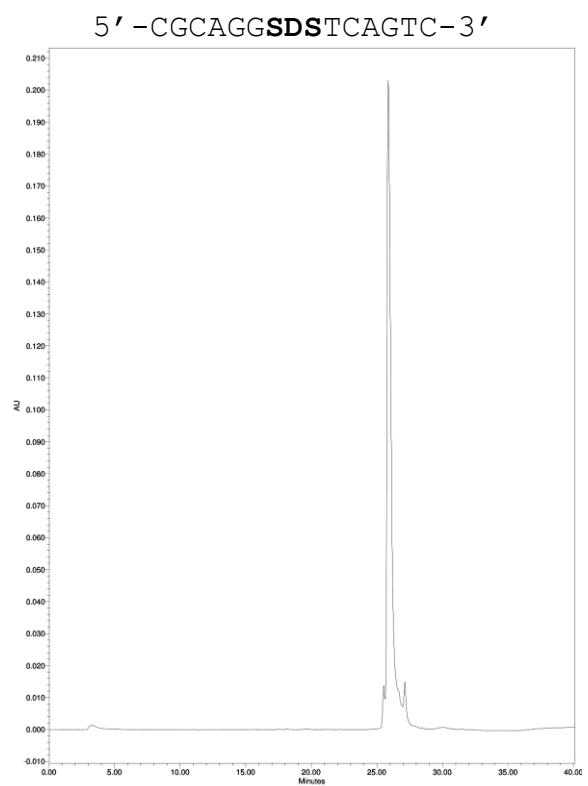
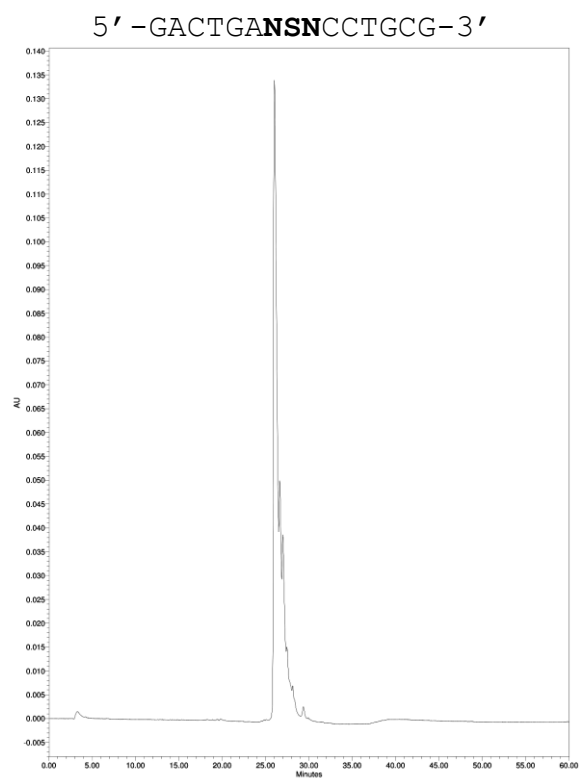


Table S1. HRMS-ESI of individual oligonucleotides.

Sequence	Ion	Mass (calcd)	Mass (found)
5'-GACTGACCTGCG-3'	[M-4H] ⁻⁴	910.1545	910.1540
5'-CGCAGGTCAGTC-3'	[M-4H] ⁻⁴	910.1545	910.1537
5'-GACTGAAAACCTGCG-3'	[M-5H] ⁻⁵	915.7567	915.7570
5'-CGCAGGTTTTTCAGTC-3'	[M-3H] ⁻³	1517.9211	1517.9144
5'-GACTGAGCGCCTGCG-3'	[M-5H] ⁻⁵	917.3524	917.3528
5'-CGCAGGCGCTCAGTC-3'	[M-3H] ⁻³	1516.2568	1516.2541
5'-GACTGASSSCCTGCG-3'	[M-4H] ⁻⁴	1013.6607	1013.6586
5'-CGCAGGSSSTCAGTC-3'	[M-5H] ⁻⁵	810.7271	810.7270
5'-GACTGADSDCCTGCG-3'	[M-3H] ⁻³	1505.2795	1505.2785
5'-CGCAGGSNSTCAGTC-3'	[M-5H] ⁻⁵	875.1389	875.1373
5'-GACTGANSNCCTGCG-3'	[M-5H] ⁻⁵	939.5506	939.5497
5'-CGCAGGSDSTCAGTC-3'	[M-4H] ⁻⁴	1071.1842	1071.1818

Figure S27. HPLC Chromatograms of oligonucleotides made with 1, 2, and/or 3.





Method for Determining Extinction Coefficient of **1** and **2**

The extinction coefficient of **1** and **2** were determined based on dilutions and UV measurements of water-soluble DAN and NDI analogues (**Figure S28**). The DAN and NDI analogues were diluted with a phosphate buffer (pH 7, 100 mM NaCl, 10 mM NaH₂PO₄, 0.1 mM EDTA) and UV absorbance measurements were taken at 260 nm. The resulting extinction coefficients (2504 and 1955 M⁻¹•cm⁻¹) were used as an estimate for the corresponding phosphoramidites **1** and **2**, respectively. Because the content of the non-natural DNA base analogues is small compared to the amount of naturally occurring DNA bases, this approximation is unlikely to cause any major discrepancies in DNA concentration.

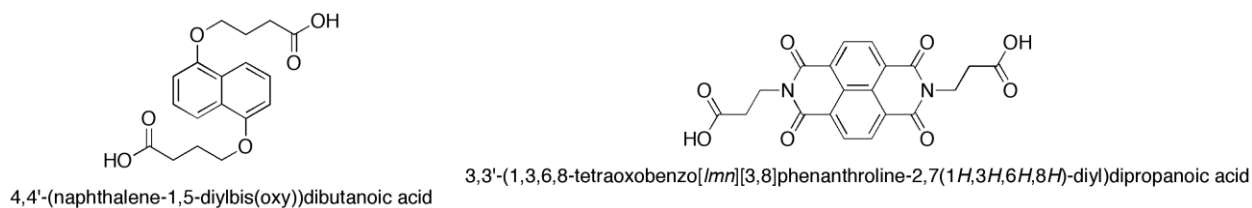


Figure S28. Structures of DAN and NDI analogues used to determine extinction coefficients for **1** and **2**.

Figure S29. UV melting curves of DNA duplexes.

